SEO ID	SEO ID NO:	Met	ISEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
İ	sequence	į	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1				sequence		
21775	52143	A	21903	843	2055	HHRMAQNLSRYGNDKRPEESQ
		1		1		FRVVNVKPVTLYDVAEYAGVS
				1		YOTVSRVVNOASHVS\AKTREK
						VEAAMAELNYIPNRVAQQLAG
	1	1				KQSLLIGVATSSLALH\APSQIV
1	ł	1			l	AAIKSRADQLGASVVVSMVF\R
l						SGLVACHTAPHNLL/ARKRGPA
		1				GLIINYPLDD\QDAIAVEA\ACT
1		i			l .	NVPALFLDVSDQTPINSIIFSHE
					i	DGTRLGVEHL\VALGH\OOI\AL
1		1				LAGPLSSVS\ARLRLAG\WHNFL
1	1	1				TGN\QIQPIA\ER\EGDWSAMSGE
1		1				QQT\LQMLNEGIVPHWRLLVAQ
ŀ		ŀ		l		RSRMALG\AMRRHYPESGLRV
	ŀ	1				GA\DILRVGGNTTIPED\SSCYIP
		1		ŀ		AL*PPIKQDFRLLGA/QTSVDR\L
		1			l	LOLSOGOAVKGN\QLLPVSL\V
1				1		KRKTTL\APNTQTA\SPRALPRF
01776	50144	١.	21004		1201	H*MQLA\RQVSRLESGQ
21776	52144	A	21904	1	1201	
21778	52146	A	21905	646	934	
21779	52140	A	21900	1	1713	MLCTKWDKLGLEKNAGKNKN
21//9	32147	^	21907	l '	1773	AYKWYGWPANAPDEPHIEOLV
1	Į.	ĺ				GECRVMROLKRLISRIAPSPSSV
1		1				MVVGESGTAIPVVSHAEFKGGF
			l			ADIGVHYLDWTSRTTEKSSTKS
		1				HKDDFGYLEFEGGANFSWGEM
1		1				YGFFDWENFYNGRHNKPGSEQ
		1				RYTFKNTKPYLSWVDTGFSLYL
1		1				HAYGTYGSANRVNFLDDM\FL
						AGFGFQF*PAGGW/WGSNPFFA
						K\RYTRSKPYYT\GDNGYV\AG
		1				WVAGYNFMLGSEKFTLTNWN
		1				EYEFDRDATYAADFLPLYDVD
						CODNGNLEYDTYSOPEWKHNL
						FDHYLAVLYRFKDESGKEQFSC
						AVVKTREATPGKEIEAITRRML
						DFSPRLKKLADCPSPRPVFEALO
1		1	1		1	ICSMLDALLLLCPLDYHASCDN
		1				YSYALLDPQGFPLPHRRLWGPY
		1	1	1		
		1	1		1	IDVDVPMLHLIQHAWAQDALL
	1	1				SCLYSDLLYMSSFPYPAGTLTR
		1	1	1	1	LMGVTSYNHKNTLLTYCITLMS
		1	1	1	1	TIRFAYDGDEEIAMKYTDFLKG
	1	1	1	1	1	ELSMNIRPLHDRVIVKRKEVET
		1	1		1	KSAGGIVLTGSASAKSTRGEVL
	I	1	1	1	1	AVGNGRILENGEVKPLDVKIGD
L		<u> </u>				IVFFNDGYRGEI
21780	52148	Α	21908	2060	2404	
21781	52149	В	21909	1	1377	

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		_				
21782	52150	Α	21910	52	1719	IGQVVDVDTCFTRIHLVFVNFD
				1		NDTGSIHVLNDTTTFSYRSYTG
		ŀ		i		VNGNSTFHTSTNQRLISAQSRN
						GLTLHVRTHQCTVGVIVFQERD
		ļ.		Į.		QGRTDGYHLLGGYVHVVNLV
i i		l				AAEQAGFAFATASYQVFYEVA
				ĺ		FFIQVGVRLGDNVVAFFDSRQV
						VNFVSYNTVGHFTIRSLKEAVF
						VSLCVHGQRVDQTDVRTFRGF
		1				DWTDATVVSRVYVSNFEACTF
						TGQTAWAECGDTTFVRNLRQR
		1				VVLVHKLRQLAGTEELFHCCG
						SQDPQQPWGWNLRVNPLAQTT
						GHAHEAQRKQWHKGNEEADD
						PEPERAFTPGFVQFETERFRPPV
						GHPCEAAEDHAADDHVMEVR
						NQEQAVMQNEVRARYGQQHA
		1		l		SHPAHREGDDKTDGPQHRRME
1						HDAALIHGEQPVEDLHPGRDG
ŀ						DNHCGDTKEGVYVRARTHRKE
l						VVQPNDERQHGDTDGCPYQRG
		1				ITKQTFTREGCRNFGEHTENRQ
		1		l		DQDVNFRVAPGPNQVDVHHH
						VATHIVGKEMGAQIAIQGQHAT
				i		YRQNGKDL*SNRGVASGNPQC
		ŀ				RHRAHRRATDCOTVIKOPVFFG

	SEQ ID NO:					Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21783	52151	Α	21911	608	2508	YITATSCSGWMIRWCAKAAKC
						CA/CSRGPGGKLRRHKGAEETL
						CGNTRVLIAPQLPAVIIATLSGS
						RSQTGRREGPTPTTVMIQDHTG
						SSTDDECSANVQRIFNRIMLCSI
						VSRPRTIPFHDLAQSFRQFRNVK
						CLWCAFEGAQLMRIQPAIVMV
						KTIHRERLNSGQFRCHQEARNI
						PPHRQSRFLASRGLLAELMFML
			İ			YGIGELPEIVTLPKEIRGNVDNS
						RHAVFLANSGMPELRFIRTHGS
						RFTSQDCTLFNWLARIITPVLQS
ł		l				WLNDEEQQVALRLLEKDRDHH
						RVLVDITNAVLSHLDLDDLIAD
		1		1		VAREIHHFFGLASVSMVLGDHR
		1				KNEKFSLWCSDLSASHCACLPR
1						CMPGESVLLTQTLQTRQPTLTH
		ļ				RADDLFLWQRDPLLLLLASNG
			ŀ			CESALLIPLTFGNHTPGALLLAH
		l				TSSTLFSEENCQLLQHIADRIAI
			İ			AVGNADAWRSMTDLQESLQQ
	1					ENHQLSEQLLSNLGIGDIIYQSQ
	1	l				AMEDLLQQVDIVAKSDSTVLIC
1	l					GETGTGKEVIARAIHQLSPRRD
	1					KPLVKINCAAIPASLLESELFGH
	1					DKGAFTGAINTHRGRFEIADGG
	İ	1				TLFLDEIGDLPLELQPKLLRVLQ
						EREIERLGGSRTIPVNVRVIAAT
		_				NRDLWQMVEDRGVI
21784	52152	A	21912	147	2699	MTIRAICSCCPLPDA*NVLM*IY
		l				MSTFQASGNGQQLQIARIVIQNI
		L.				TG
21785	52153	Α	21913	432	1016	FCPSSCLAATAPAA*AGWSFLP
	l					SPAEQNRWRFPLSRFCQIPAPL/
	1	1				LGTLIISLSVFCSILAASAAEIPG
1	1	l				SVDGIYNKSPSLI\WVKIPRQLY
1	1	l				*AAREPSAPKAPPPAAWFSASA
		1				VLSATA\MINTNQKAIERVTAFR
				1		RNFATDPVAHQDRDHYDRQQR
	1	l				RTRHRIRFGERQRAKQTPFLPFQ
		<u></u>				GEDRDKRQRDNQQTDK

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	1			sequence		
		_				
21786	52154	A	21914	I	1146	MTTSMLNAKLLPSAPSAAVVV
						VRVVVVVGNAPTSRKVGPEEK
						DWSMASSGTTSTRKRFTGAEFI
						VHFLEQ/PGH*DCDRHSG/S/CSI
	1					LPVYDALSQSTQIRHILARHEQ
						GAGFIAQGMARTDGKPAVCMA
1						CSGPGATNLVTAIADARLDSIPL
						ICITGQVPASMIGTDAFQEVDT
		1				YGISIPITKHNYLVRHIEELPQV
						MSDAFRIAQSGRPGPVWIDIPK
1				l		DEQTAVFEIETQPAMAEKAAAP
						AFSEESIRDAAAMINAAKRPVL
						YLGGGVINANARVRELAEKAQ
						LPTTMTLMALGMLPKAHPLSL
ľ						GMLGMHGVRSTNYILQEADLLI
		1				VLELAGVOTVDTDVNASKPAS
		1				ATRHIAFOTVTIGCDRNLADSV
i						VFPHSVDDIGKVAA
21787	52155	A	21915	1001	1037	MVGMASSCMPAFTRLELYSVIS
21/0/	32133	^	21713	1001	1037	QSTQIRH/ILARHEQGAGFIAQG
						MA/RTDGKPAVCMACSGPG/AT
						NLVTAIADARLDSIP/LICITGOV
1		l				PASMIGTDA/FOEVDTYGI/STM
		l	1	l		TLMALGMLPKAH/ALSLGMLG
			i		1	
		İ				MHGVRSTNYFCQEADLSIVLG
						ALFDAGQIGKTSSFCPNAKI/IH
			l			VDIDRAE/REFPCPIPKACDLLS
1	ļ					H/YGLINAVAACVDDNAIITTD
	1	l	1			VGQHQMWTAQAYPLNRPRQW
			1			LTSGGLGTMGFGLPAAIGAALA
	1		1			NPDRKVLCFSGDGSLMMNIQ*
						MIIAIPTGVKIFNWCSPCIRPHR
						VPFCDAVAIGFIVTFSDVIQIASI
						YEIPCRDPTGVASTEYSSSSCRR
						RRNTTLCDPGSPAQRQSPQAGQ
		1_				NPSCPAHRRSATGVGD
21788	52156	Α	21916	135	449	
21789	52157	Α	21917	5	214	
21790	52158	A	21918	2	337	IGIAKHAQQLLLNVMQTRAVG
		1				AEVQTVSATGSAVRQALANNL
		1		l		QPFAIFFQINA/W**RAGRYPDD
1	1	1	1	I	1	HGISPETALYGWHTSATGTAG
		l		1	[MR*SVRPWLQSGQGCSGWRRQ
1		1	1			OPRWSP
21791	52159	A	21919	1	774	-
	1	11.				J

| SEQ ID NOC. Net SEQ ID NOC. Nocleoside Noc. Nucleoside Noc. of peptide NVADG
LRADS
KNTPL
PLAVW
KRLV
NAAQ
VTSV*
ALRAG
DRQSV |
|--|---|
| 21792 52160 A 21920 2 914 RHIRTPAHLERFIVDIALL HRAKLIVOSAGITRAQITV SANNERGOVGLVROPERE VARQAAVSLARFICLAF VARQAAVSLARFICLAF VARQAAVSLARFICLAF NFNKLNFADLQRFLWRI LSCAITIT/ADPAVLLHH PAQI-WRIS-PYRONLET. MI-HMI.GDKTT/MI-FIRM NA*DLOPFRVNAFAEIA VSETAGHARTEVNAGFI DTTRHIFFITVANAFAYH VTHAEFITRAACGIQITT QAGVADNACEMAAEG TNGD | NVADG
LRADS
KNTPL
PLAVW
KRLV
NAAQ
VTSV*
NLRAG
DRQSV |
| 21792 52160 A 21920 2 914 RHIRTPAHLERFIVDIALI HRAKLIVQSAGTFAQTV SANFRQGVGLVRQFRRF VGELQPVRDVYVRAFI VAARQAAVSLRFGLAFG NFNKLNFADLQRFLWRI LSCATTT/ADPAVLLHH PAQI*WRIS*PVRQNLET: MLHMLGDKTTQMLFM NA*DLDQFRVNAFAEIA VSETAGHARIEVNAGFI DTTRHIFTIVVANAFVH VTHAETFRAAGGIQTT QAGVADNACFMAEGG TNGD | LRADS
KNTPL
PLAVW
KRLV
NAAQ
VTSV*
ALRAG
DRQSV |
| HRAKLIVQSAGTFAQTV SANFRQRYGLVRQFRR VGELQPVRDVVYYRAFI VAARQAAVSLRFGLAF* VAARQAAVSLRFGLAF* NFRIKLIFADLQRFLWRI LSCATTIADPAVLLHH PAQI*WRS*PYRQNLFT MLHMLGDKTTQMLFMI NA*DLDQFRVNAFAELA VSETAGHARIEVNAGFI DTTRHIFTIVVANAFVH VTHAEFTERAAGIQTY QAGVADNACFMAAEGI TINGD 21793 52161 A 21921 204 371 PGPAGAQSAALYRWSPC RADWFRADC*IYPERA AYRRLSPAYGI AYRRLSPAYGI A 21922 2664 2970 TPDTAPAGCSSRAPDGP MPRRG*FHTARYCRTP *R*TE*STCPARTGEWY, APGTSRQTATMPPTMFR YSSTQPDGRRQNC 21795 52163 A 21923 318 459 21796 52164 A 21924 577 888 TNSNQQLAEGNRERLIG GPERITAGRR*LFSCQNR RFNCCHIGMPVVVIWERI QHDDLVNPFRQNEVN | LRADS
KNTPL
PLAVW
KRLV
NAAQ
VTSV*
ALRAG
DRQSV |
| HRAKLIVQSAGTFAQTV SANFRQGVGLVRQFRR VGELQPVRDVVYYRAFI VAARQAAVSLRFGLAF* VAARQAAVSLRFGLAF* VFRILNFADLQFELWRI LSCATTTADPAVLLHH PAQI*WRIS*PVRQNLFT MLHMLGDKTTQMLFM NA*DLDQFRVNAFAELA VSETAGHARIEVNAGFI DTTRHIFTITVANAFYH VTHAEFTERAAGIQTT QAGVADNACFMAAEG TNGD PGPAGAQSAALYRWSP RADWFRADC*IYPERA AYRRLSPAYG AYRRLSPAY | LRADS
KNTPL
PLAVW
KRLV
NAAQ
VTSV*
ALRAG
DRQSV |
| SANFRQGVGLVRQPRET VGELQPVRDVVYYRAF VGARQAAVSLRFGLAFC VGARQAAVSLRFGLAFC VGARQAAVSLRFGLAFC VGARQAAVSLRFGLAFC VGARQAAVSLRFGLAFC VGARQAAVSLRFGLAFC VGARQAAVSLRFGLAFC VFRIKTING LSCATTTADPAVLIHH LSCATTTADPAVLIHH LSCATTTADPAVLIHH LSCATTTADPAVLIHH LSCATTTADPAVLIHH LSCATTTADPAVLIHH LSCATTTADPAVLIHH VGAGVADVALFT VGAGVADVACFMAFGT VGAGVADVACFMAFGT VGAGVADVACFMAFGT VGAGVADVACFMAFGT VGAGVADVACFMAFGT VGAGVADVALFT VGAGVADVATAT VGAGVADVATAT VGAGVADVATAT VGAGVATAT VGAGVATAT VGAGVATAT VGAGVATAT VGAGVAT VG | KNTPL
PLAVW
KRLV
NAAQ
VTSV*
ALRAG
DRQSV |
| VGELQPYRDVVVYRAF VAARQAAVSLRFGLAF(NFNKLNFADLQRFLWRI LSCATTT/ADPAVLLHH PAQI*WRS*PYRQNLFT. MLHMLGDKTYQMLFMI NA*DLDQFRVNAFAEIA VSETAGHARTEVNAGFI DTTRHIFTTVANAFYH VTHAEFTFRAACGIQTT QAGVADNACFMAAEGC TNGD QAGVADNACFMAAEGC TNGD PGPAGAQSAALYRWSPC RADWFRFADC*1YPENR AYRPRLSPAYCG A 21922 2664 2970 TPDTA*PAGCSSRA*PDG*P MFRRG*PHTARYCST*P *R*TE*STCPARTGEWY/ APGTSRQTATMP*TMFR YSSTQP*DGRQNC 21795 52163 A 21923 318 459 21796 52164 A 21924 577 888 TNSSNQQLAEGNRERLI GPERITAGR**LFSCQNR RFNCKHIGMP*VVVIWRI QHDDLVNFRGMRVNRV | PLAVW
KRLV
NAAQ
VTSV*
ALRAG
DRQSV |
| VAARQAVSLRFGLAFF NFNKLNFADLQRFLWR LSCATTT/ADPAVLLHH LSCATTT/ADPAVLLHH PAQI*WRS*PYRQNLET: MLHMLGD&TTYDMLFM NA*DLDQFRVNAFAEIA VSETAGHARIEVNAGFI DTTRHFITIVANAFYH VTHAETFTRAACG QTT QAGVADNACFMAEGG TNGD | KRLV
NAAQ
VTSV*
ALRAG
DRQSV |
| NFNKLINFADLQRFLWRI LSCATIT/ADPAVLLHH | NAAQ
VTSV*
ALRAG
DRQSV |
| LSCATT/ADPA/LLHH | VTSV*
ALRAG
DRQSV |
| PAQI*WRIS*PVRQNLFT. MLHMLGDKTTQMLFM | ALRAG
DRQSV |
| MI.HMLGDKTTOMLFMI NA*DLDQFRVNAFAEIA VSETAGHARTEVNAGFI DTTRHIFTTVVANAFVH VTHAEFTERAAGIQTT QAGVADNACFMAAEGG TNGD QAGVADNACFMAAEGG TNGD PGPAGAQSAALYRWSPC RADWFRFADC*IYPENR AYPRI.SPAYCG RADWFRADC*IYPENR AYPRI.SPAYCG PFDTAPAGCSSRAPDGP MPRRG*PHTARYCSTP *R*TE*STCPARTGEWY/ APGTSRQTATMPPTMFR YSTCPPGRGNC VSTCPPGRGNC VSTC | RQSV |
| NA*DLOOFRVNAFAELA VSETAGHARTEVNAGFT | |
| VSETAGHARTEVNAGT | CHEN |
| DTTRHIFTTVVANAFYH | ATIEN |
| VTHAETFTRAACGIQTT QAGVADNACFMAAEG TNGD | QNAD |
| QAGVADNACFMAAEGC | SDSTG |
| QAGVADNACFMAAEGC | GRAI |
| TINGD | |
| 21793 52161 A 21921 204 371 PGPAGAQSAALYRWPPR RADWFRFADC*I/PENR AYRRLSPAYCG PFDTAPAGCRSRAPDGP PFDTAPAGCRSRAPDGP PFDTAPAGCRSRAPDGP PFDTAPAGCRSRAPDGP PFPTAPAGCRSRAPDGP PFPTAPAGCRSRAP | |
| RADWFRFADC* YPENR AYRPRLSPAYCG | OAVS |
| AYRPRLSPAYCG | |
| 21794 52162 | 4.7100 |
| MPRRG-9FHTARYCRTP *R*TE*STCPARTGEWY, APGTSRQTATHEPMY PMPTWR YSSTQPDGRRQNC | SI PRO |
| *R*TE*STC*ARTGEWY, APGTSRQTATMPPTMP YSSTQPDGRQNC 21795 52163 A 21923 318 459 21796 52164 A 21924 577 888 TNSSNQQLAEGNRERLIS GPERITAGR*LFSCQNR RFNCCHIGMPVVVIWAII QHDDLVNPRMVNRV | |
| APGTSRQTATMPPTMPR YSSTQPDGRRQNC 21795 52163 A 21923 318 459 21796 32164 A 21924 577 888 TNSSNQQLAEGNRERLIG GPERITAGRR*LESCQNB RFNKCHIGMPVVVIWRI QHDDLVNPFGMRVNRV | |
| YSSTQPDGRRQNC | |
| 21795 52163 A 21923 318 459 | idaiss |
| 21796 52164 A 21924 577 888 TINSNQQLAEGNRERLIG
 GPERITAGRR*LPSCQNIG
 RFNKCHIGMPVVVIWRI
 QHDDLVNPFGMRVNRV | |
| GPERITÄGRR*LFSCQNR
RFNKCHIGMPV-VIVIII
QHDDLVNPFGMRVNRV | RNTA |
| RFNKCHIGMPVVVIWRI
QHDDLVNPFGMRVNRV | |
| QHDDLVNPFGMRVNRV | |
| | |
| | |
| 21797 52165 A 21925 3 711 AGAHSGPSHPALATTAA | |
| 21/9/ 32163 A 2192 S MI AGNRSGISGFSGHSRNPI | |
| RCL*SS\ALERLMRRLNC | |
| SDVTTTTPS/LSPOVYCS. | |
| LCTRATPRWQSAGHPYC | |
| | |
| SYCAIYSDRSIRGNYSTR | |
| GRRV\PTGSWQKTESEK | |
| CANWANDRTPLPTSTSS | |
| NVQTSVVPVLRRWIWLT | ANCST |
| VLTVSLKSVCARGTSPQ | ANCST
SLRV |
| VKRAASSATSPVVITTC | ANCST
SLRV |
| 21798 52166 A 21926 244 535 TRYPDDHGISPETALYG | ANCST
SLRV
ASCWF |
| TGTAGMR*SVRPWLQSG | ANCST
SLRV
ASCWF |
| GWRRQQPRWSP**RSC* | ANCST
SLRV
ASCWF
WHTSA
GQGCS |
| PATL*FCRLYNQTSGPGi | ANCST
SLRV
ASCWF
WHTSA
GQGCS
CSVY |
| AFPDSDRRKI | ANCST
SLRV
ASCWF
WHTSA
GQGCS
CSVY |

SEQ ID NO:	SEQ ID NO: of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21799	52167	A	21927	869	2923	PKKDLKIATSAMMEAAYSV*SR TLFW/SPVRDSQVTDTQWQGL MSQLRLQGFDTLVLQWTRYGD AFTQPEQRTLLFKRAAAAQQA GLKLIVGLNADPEFFMHQKQSS AALESYLNRLLAADLQQA RLW SAAPGITPDGWYISAEIDDLNW RSEAARQPLLTWLNNAQRLISD VSAKPYYISSFFAGMMSPDGYR QLLEHWKATGVNVWYQDGSG VDKLTAEQRERYLQASADCHD VDKLTAEQRERYLQASADCHD
21800	52168	В	21928	72	690	
21801	52169	A	21929	1	2241	
21802	52170	A	21930	1607	1893	NKRIFPSDVKVAEPDHSCV*SFR IGSLASNLSLSLKTRPAFFQLLIK HVD/SMCHQRLSGSRKSKHNVI AVVFSYCSVSVFSASCPTHQVL LQTQYF
21803	52171	Α	21931	1	3960	
21804	52172	Α	21932	1	1281	
21805	52173	Α	21933	1	1491	
21806	52174	Α	21934	1	1254	
21807	52175	Α	21935	1	723	
21808	52176		21936	740	880	NKRIFPSDVKVAEPDHSCV*SFR IGSLASNLSLSLKTRPGFFPTVN
21809	52177		21937	1	1191	
21810	52178	Α	21938	1033	1248	
21811	52179	Α	21939	1	1425	
21812	52180	Α	21940	1	2784	
21813	52181	Α	21941	676	1467	
21814	52182	В	21942	1	3288	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /-possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21815	52183	A	21943	145	1929	VSGVIEIADGSRRRKAAALTES
21013	32163	<u>۱</u> ^	21943	143	1929	
						DYRVLVGELDDEQMAALSRLG
						NDYRPTSAYERGQRYASRLQN
						EFAGNISALADAECDNLKTCHT
		l		}		SHGSVMAETAVINHKKRKNSP
				ł		RIVQSNDLTEAAYSLSRDQKRM
		ŀ		i		LYLFVDQIRKSDGTLQEHDGIC
		1				EIHVAKYAEIFGLTSAEASKDIR
		l				QALKSFAGKEVVFYRPEEDAG
		l				DEKGYESFPWFIK/RPSRGLYSV
ı		l				HINPYLIPFFIG\LQNRFTQFRLS
i						ETKEITNPYAMRLYESLCQYRK
	1					PDGSGIVSLKIDWIIERYQLPQS
i						YQRMPDFRRRFLQGFCRFRNH
l	1	1				HQTGFSPAGANQRGPLAATLSG
i		1				PGGEGQSAVARLTGEKKNHPG
i						AQYANRLSPRVGRFINAAGTTG
i						FPTWKAGSERNAINDDVTYAIK
i						PTCWPGLDIIPSCLALHRIETEL
i		l				MGKFDEGKLPTDPHLMLRLAIE
i						TVAHDYDVIVIDSAPNLGIGTIN
i						VVCAADVLIVPTPAELFDYTSA
i						LQFFDMLRDLLKNVDLKGNSN
l		1				GSQSPWMEEQIRDAWGSMVLK
i						NVVRETDEVGKGQIRMRTVFE
i						QAIDQRSSTGAWRNALSIWEPE
						CNEISIGVSLDQDGGSNSVLRK
21816	52184	Α	21944	1790	2947	TDFSVRCKGCRTGPFLRLIFQDR
ı						IVGVELIAVF*NTGPRFSSCRLS
i .						WRRPFDKTCRMRWHQRLSGFP
						QEQNDGKQQPILC
21817	52185	Α	21945	1	888	
21818	52186	Α	21946	252	1863	RKNPFILH*LFR*TLRQTKPDDS
						AGKGVKI**HTQNQRSGRSQND
						F*RRLW
21819	52187	В	21947	1	1794	

NO. of peptide requence hod with 28/31/17 in USSN with reduction of first codons for patide sequence codon for patide sequence codon for peptide deletion, "possible nucleotide insertion) 21820 32188 A 21948 138 341 PNPRGLLNTITPEHFKSKLHIPT TYRQSNGITASHYSYARN*R*N.W.WARGNAITLPVCGRDVKFTI EVLRGDSVEKTSRVWSGLLEL KAAASLVFLEOHEYPEEIGTS GSVLVSVVGEOPFVEEDLRS. PGRWDKKHKMFDAIEPPKKD: DKFDANDPILKDQTGEWSGSA TFTSDGKIRLFYTDYSGKHYGI QSLTTAQVNVSKSDDTLKING EDHKTIPEDGGGKTYQNVQQFI EGNYTSGDNHTLRDPHYVEED GHKYVFEANTGTENOYQCGI SLFNKAYYGGGTNFFRKESQK QQSAKRRDALANDAGIEIEL YDYTLKKMKAADHFQTPVT EIERANVFKNMG 21821 32189 A 21949 761 916 NKRIFPSHVKVAEPDHSCV*SF IGSLASNLSLSLKTRPAFFQLSI LSPT 21822 32191 B 21951 50 1690 1437 21823 32191 B 21951 1002 257KKSLASNLSLSLKTRPAFFQLSI LSPT 21826 32193 A 21954 878 1202 SVYNKRIFPSDVKVAEPDHSCV *SFRIGSLASNLSLSLKTRPAFF 21826 32194 A 21955 7 1702 VSFRIGSLASNLSLSLKTRPAFF 21827 32195 B 21955 7 1702 NKRIFPSDVKVAEPDHSCV *SF GOS 21829	SEO ID	ISEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
21820							
21820 52188 A 21948 138 341 PNPRGLLNTITPEHFKSKLHIPT TYRQSNGITASHYSYARN*RViv VMARGNAITLPVCGRDVKFT1 EVLRGDSVEKTSRVWSGLLEL KAAAASLVFLEGHEYPEGEGIST GSVLVSVVGEGPFVEEDLRKS PGRWDKKHKMFDAIEEPKKD: DKFDANDPILKDQTGEWSGSA TFTSDGRIRLFYTDYSGKHYGI QSLTTAQVNVSKSDDTLKING EDHKTIPEDGGKTYQNVQQFI EGNYTSGDNHTLRDPHYVEDG GHKYLVFEANTGTENGYQGEI SLFNKAYYGGGTNFFRKESQK QQSAKRRDAELANGALGIELI YDYTLKKVMKAADHFQTPVT EIERANVFKMNG NKRIFPSHVKVAEPDHSCV*SF IGSLASNLSLSLKTRPAFFQLSI LSPT 1437 21822 32190 A 21950 I 1437 21823 32191 B 21951 30 1690 21824 32192 B 21952 I 1032 21825 32193 B 21953 I 1974 21826 32194 A 21954 878 1202 SVYNKRIFPSDVKVAEPDHSCV*SF LSIERFRPFFCKELMCHQRL; GFPHEQTQRDSTCLQLFRQRF AAASCPTHQVLLQTQYF 21827 32195 B 21955 7 1702 7 1702 7 1702 7 1702 1					codon for peptide		deletion, \=possible nucleotide insertion)
TYRQSNGITASHYSYARN*RV VMARGNATILPYCGROWEFTE EVLRGDSVEKTERVWSGLLE KAAASLYFLEQHEPYEEJGTSS GSVUSVVGEPPYEEJGTSS GSVUSVVGEPPYEEJGTSS GSVUSVVGEPPYEEJGTSS GSVUSVVGEPPYEEJGTSS GSVUSVVGEPPYEEJGTSS GSVUSVVGEPPYEEJGTSS GSVUSVVGEPPYEEJGTSS GSVUSVVGEPPYEEJGTSS GSVUSVVGEPPYEEJGTSS GSLATAQVNVSSSDDTLKING GEHKTIPEDGBKTYQNVQOFF ECNYTSGDHTLRDPHYVEDG GHKYLVFEANTGENGYQGE ECNYTSGDHTLRDPHYVEDG GHKYLVFEANTGENGYQGE SLFNKAYYGGGTNFFKESQK QQSAKKROAELANGALGIELI YDYTLKKVMKAADHFQTPYT EIERANVFKMNG NKRIPPSHVKVAEPPHSCV*SF IGSLASNLSLSLKTRPAFFQLSI LSPT					sequence	1	
TYRQSNGITASHYSYARN*RV VMARGNATILPYCGROWEFTE EVLRGDSVEKTERVWSGLLE KAAASLYFLEQHEPYEEJGTSS GSVUSVVGEPPYEEJGTSS GSVUSVVGEPPYEEJGTSS GSVUSVVGEPPYEEJGTSS GSVUSVVGEPPYEEJGTSS GSVUSVVGEPPYEEJGTSS GSVUSVVGEPPYEEJGTSS GSVUSVVGEPPYEEJGTSS GSVUSVVGEPPYEEJGTSS GSVUSVVGEPPYEEJGTSS GSLATAQVNVSSSDDTLKING GEHKTIPEDGBKTYQNVQOFF ECNYTSGDHTLRDPHYVEDG GHKYLVFEANTGENGYQGE ECNYTSGDHTLRDPHYVEDG GHKYLVFEANTGENGYQGE SLFNKAYYGGGTNFFKESQK QQSAKKROAELANGALGIELI YDYTLKKVMKAADHFQTPYT EIERANVFKMNG NKRIPPSHVKVAEPPHSCV*SF IGSLASNLSLSLKTRPAFFQLSI LSPT			<u> </u>				
VMARGHAITLPVGGROVEFTE EVLRGDSVEKTSRVWSGLEL KAAASLVFLEQHEPVEEIGTS GSVLVSVVGEGPFVEEDLRS PORWOKHIKRIPDAIEPKEND DKFDANDPILKDOTGEWSGSA TFTSDGKIRLFYTDYSGKHYGI QSLTTAQVNVSKSDDTLKING EDHKTIFDGDGKTYQNVQQFI EGNYTSGDMTHLRDPHYVED GHKYLVFEANTGTENGYQGEI SLFNKAYYGGGTNFFRKESQK QQSAKKRDAELANGALGIIELI YDYTLKKVMKAADHFQTPVT EIERANVFKNNG NKRIFPSHVKVAEPDHSCV*SF IGSLASNLSLSLKTRPAFFQLSI LSFT	21820	52188	Α	21948	138	341	
EVLRGBSVEKTSRVWSGLEL KAAASLVFLEQHKPYEEIGTSI GSVLVSVVGEOFPVEEDLRKS PORWDKKHKMFDAIEPFKDD DKFDANDPILKDOTQEWSGSA TFTSDGIRLFYTDVSGKHYGI QSLTTAQVNVSKSDDTLKING QSLTTAQVNVSKSDDTLKING GPHKTIPEDGDKTYQNVQOFF ECMYTSGDNHTLRDPHYVEDG GHKYLVFEANTGTENGYQCE SLFNKAYYGGGTFFRKESQK QQSAKKDAELANGALGIIELI YDVTLKKVMKAADHFQTPVT EIRBANVFKING POTTLKVMKAADHFQTPVT EIRBANVFKING EIRBANVFKING CSLASNLSLSLKTRPAFFQLSI LSPT LSP					İ		TYRQSNGITASHYSYARN*RVG
KAAASLYPLEQHRPYEEIGTSK GSVLVSVVGEGPFVEEIDLRKS PORWDKKHKMFDAIEPPKKDD PORWDKKHKMFDAIEPPKKDD PORWDKKHKMFDAIEPPKKDD PORWDKHKMFDAIPIKKDQ QSLTTAQVNVSKSDDTLKING QSLTTAQVNVSKSDDTLKING EDHKTIPDGDGKTYQNVQQFI EGNYTSGDNHITLRDPHYVEDB GHKYL-VFEANTGTENGYQGE SLFNKAYYGGGTFFRKESGK QQSAKKRDAELANGALGIELI YDYTLKKVMKAADHFQTPVT EIERANVFKNNG NKRIFPSHVKVAEPDHSCV*SF USLASNLSLSLKTRPAFFQLSI LSPT SIGSLASNLSLSLKTRPAFFQLSI LSPT SIGSLASNLSLSLKTRPAFFQLSI LSPT SIGSLASNLSLSLKTRPAFFQLSI LSPT SIGSLASNLSLSLKTRPAFFQLSI LSPT SIGSLASNLSLSLKTRPAFFQLSI SIGSLASNLSLSJKTRPAFFQLSI SIGSLASNLSLSLKTRPAFFQLSI SIGSLASNLSLSLKTPAFFQLSI SIGSLASNLSLSLKTPAFFQLSI SIGSLASNLSLSLKTPAFFQLSI SIGSLASNLSLLKTPAFFQLSI SIGSLASNLSLLKTPAFFQLSI SIGSLASNLSLLKTPAFFQLSI SIGSLASNLSLLKTPAFFQLSI SIGSLASNLSLLKTPAFFQLSI SIGSLASNLSLKTPAFFQLSI SIGSLASNLSLASNLS SIGSLASNLS		1	1				VMARGNAITLPVCGRDVKFTL
GSVLVSVVGEGPFVEEDLRKS PGRWDKKHKMFDAIEEPKKD: DKFDANDPILKDQTQEWSGSA TFTSDGRIRLFYTDYSGKHKGH GVSLTTAQVNVSKSDDTLKING GVSLTTAQVNVSKSDDTLKING GUSLTTAQVNVSKSDDTLKING GUSLTAQVNVSKSDDTLKING GHKYLVFEANTGTENGYQGE SLFNKAYYGGGTNFFRKESQK QQSAKRDAELANGALGIELI YDYTLKKVMKAADHFQTPVT EIERANVFKING GISLASNLSLSLKTRPAFFQLSI LSPT S12182 S2190 A 21950 I 1437 S121823 S2191 B 21951 50 1690 S121823 S2191 B 21951 J 1032 S121824 S2192 B 21952 I 1032 S121826 S2194 A 21954 878 I 1202 SVYNKRIFPSDVKVAEPDHSCV SFRIGSLASNLSLSLKTRPAFF LSIEFREPFDKTCRLMCHQRL GFPHEQTQRDSTCIQLLFRQRF AAASCPTHQVLLQTQYF AAASCPTHQVLLQTQYF AAASCPTHQVLLQTQYF S121828 S2196 A 21956 S27 665 NKRIFPSDVKVAEPDHSCV SFRIGSLASNLSLAKTRPAFF LSIEFREPFDKTCRLMCHQRL GFPHEQTQRDSTCIQLFRQRF AAASCPTHQVLLQTQYF S121829 S2197 A 21957 469 687 GSDSRAFAKENNQKAYKETY GVSHTRHDMLQIPKQQQNEK GVSHTRHDMLQIPKQQQNEK GVSHTRHDMLQIPKQQQNEK KPDANDPILKDYAEY GVSHTRHDMLQIPKQQQNEK KPDANDPILKDYAEY GVSHTRHDMLQIPKQQQNEK KPDANDPILKDYAEY KVGDNSIDSWKNAGRYKDSI KPDANDPILKDYQTQSKSAL KPDANDPILKDYQTQSKSAL KPDANDPILKDYQTQSKSAL KPDANDPILKDYQTQSKSAL KPDANDPILKDYQTQSKSAL KPDANDPILKDYQTQSKSAL KPDANDPILKDYQTQSKSAL TTDGGRIRLFYTDYSGKHYGQ LTTAQVNVSKSDDTLKINGSK KPDANDPILKDYQTQSKSGAL KP			l				EVLRGDSVEKTSRVWSGLLELL
PGRWDKKHRMFDAIEPPKNDD						l	KAAASLVFLEQHKPYEEIGTSF
DKFDANDPILKDQTGE-WSGSA TFTSDGKIRLFYTDYSGKHYG QSLTTAQVNVSKSDDTLKING EDHKTIFDGDGKTYQNVQKFI EGNYTSGDNHTLRDPHYVED GHKYL-VFEANTGTENGYQCE SLFNKAYYGGGTNFFRKESQK QQSAKKRDALANGALGHELI YDYTLKKVMKAADHFQTPVT EIERANVFKNNG NKRIFSHVKVAEPDHSCV*SF IGSLASNLSLSLKTRPAFFQLSI LSPT			1		l		GSVLVSVVGEGPFVEEDLRKSF
TFTSDGRIRLFYTDYSGKHYGI QSLTTAQVNVSKSDDTLKING EDHKTIPDGDGKTYQNVQKFI EGNYTSGDNHTLRDPHYVEDR EGNYTSGDNHTLRDPHYVEDR GHKYLVFEANTGENGYQGE SLFNKAYYGGGTFFFRKESQK QQSAKKRDAELANGALGIIELI YDVTLKKVMKAADHFQTPVT EIERANVFKING 21821 32189			l		l		PGRWDKKHKMFDAIEEPKKDS
QSLTTAQVNVSKSDDTLKING EDHKTIPDGDGKTYQNVQQFI ECMYTSGDNITLRDPHYVEDH GHKYILVPEANTGTENGYQCEI SLFNKAYYGGGTFFRKESQK QQSAKKRDAELANGALGIELI YDVTLKKVMKAADHFQTPVT EIERANVFKNNG QYSAKKRDAELANGALGIELI YDVTLKKVMKAADHFQTPVT EIERANVFKNNG NKRIFPSHVKVAEPDHSCV*SF ISSLASNLSLSLKTRPAFFQLSI LSPT SPT ST ST ST ST ST ST			1				DKFDANDPILKDQTQEWSGSA
EDHKTIPGOGKTYQNVQQFI		l	1	. 6.			TFTSDGKIRLFYTDYSGKHYGK
EDHKTIPGOGKTYQNVQQFI			1				OSLTTAOVNVSKSDDTLKINGV
EGNYTSGONHTLRDPHYVED#		1	1				
GHKYLVPEANTGTENGYQCE SLFNKAYYGGGTNFFKESQK QQSAKKRDALANGALGHEL YDYTLKKVMKAADHFQTPVT EIERANVFKNNG 21821 32189			1				
SLFNKAYYGGGTNFFRKESQK QQSAKKRDAELANGALGIELI YDYTLKVMKAADHEQTPYT DYTLKVMKAADHEQTPYT EIERANYFKNNG			1				
21821 32189 A 21949 761 916 NKRIFPSHVKVAEPDHSCV*SF		1	1		ł	1	
21821 32189 A 21949 761 916 NKRIFPSHVKVAEPDHSCV*SF IGSLASNLSLSLKTRPAFFQLSI LSPT			1				
EIERANYFKNNG			1				
21821 52189 A 21949 761 916 NKRIFPSHVKVAEPDHSCV*SF IGSLASNLSLSKTRPAFFQLSI LSPT			1				
IGSLASNLSLSLKTRPAFFQLS 21822 32190	21821	52189	Δ	21049	761	916	
LSPT	21021	100	ľ	21,545	/01	710	
21827 S2190			1				
21823 \$2191 B 21951 \$0 1690	21922	52100	1	21050	,	1427	LSF1
21826 \$2192 B 21952 I 1032							
21825 \$2193 B 21953 1 1974							
21826 32194							
**SFRIGSLASNISLSLKTRPAFF SIEFRRPFDKTCRLMCHQRL: GPHEQTQRDSTCQLLFRQRE GPHEQTQRDSTCQLLFRQRE AASCPTHQVLLQTQYF AASCPTHQVLLQTQYF 21828 \$2196 A 21956 527 665 NKRIFPSDVKVAEPDHSCV*SF IGSLAFNLSLALKTPARVFSTV IGSLAFNLSLALKTPARVFSTV IGSLAFNLSLALKTPARVFSTV 21830 \$2198 A 21958 2 1018 EGSDSRAFAKENNQKAYKETV GVSHTTRHDMLQIPKQQONEK QVPQFDQSTIKNIESAKGLDVW DSWPLQNADGTVAEYNGYIV VFALAGSPKDADDTSIYMFVQ KVGDNSIDSWKNAGRVFKDSI KPENADPILKDOTQEWSGSAT TSDGKIRLFYTDYSGKHYGKQ LTTAQVNVSKSDDTLKINGVE HKTIFDQDRKTVQNGQCRLESPWQT, GVSPTVQKLKSLKFDVRQQE VWEKDEALANWTVPNITCD NLKTCHTLHGSVMAETAVFNP KKLKNSPRIVQSNHLI			_				SVYNKRIEPSDVKVAEPDHSCV
LSIEFERPPENTCRLMCHQRL: GFPHEQTQRDSTCQLLFRQF AAASCPTHQVLLQTQYF AAASCPTHQVLQTQYF AAASCPTHQVLQTQYF AASCPTHQVLQTQYF ABABCPTHQVLQTQYF GSLAFNLSLALKTPARVFSTV IGSLAFNLSLALKTPARVFSTV GSLAFNLSLALKTPARVFSTV GSLAFNLSLALKTPARVFSTV GVSHITHEHDMLQIPKQQQNEK QVPQFDQSTIKNIESAKGLDV DSWPLQNADGTVAETYNGTYH VFALAGSPKDADDTSIWMFVQ KVGDNSIDSWKNAGRVFKDSI KFDANDPILKDQTQEWSGSAT TSDGKIRLFYTDYSGKHYGKQ LTTAQVNVSKSDDTLKINGVE HKTIFDGDRKTVQNGQGE KVGDNSIDSWKAMATVFNORTCO ULTTAQVNVSKSDDTLKINGVE HKTIFDGDRKTVQNGQGE VWEKDEALANWTVPNITCO NUMERICAL STATES CONSTRUCTION CONSTRU	2.020	2.,,,	ľ.	21,754	0,0	1202	
GPPHEQTQRDSTCIQLLFRQRF			l				
AAASCPTHQVLLQTQYF			l				
21827 \$2195 B 21955 7 1702		1	l				
21828 32196	21927	52105	D	21055	7	1702	AAASCITIQVEEQIQII
IGSLAFNLSLALKTPARVFSTVI 21829 32197			_				NK D I E D C U A E D D H C C U * C E D
21829 \$2197	21020	2170	ľ` `	21750	J-27	003	
21830	21820	52107	Δ.	21057	460	687	IOSE/A NESE/AERTI/ARVISTVE
GVSHITRHDMLQIPKQQQNEK QVPQDSTIKNIESAKGLDVW DSWPLQNADGTVAETYNGYHY VFALAGSPKDADDTSIVMFYQ KVGDNSIDSWKNAGRVFKDSI KFDANDPILKDQTQEWSGSAT TSDGKIRLFYTDYSGKHYGKQ LTTAQVNVSKSDDTLKINGVEI HKTIFDDDRKTYQNGQFIDE NIPPATPYAERPYYFEDKGLKY LGKPTVEPSVCGGRLESPWQT, GVSPTVQKLKSLKFDVRGQEE VWEKDEALANWTYPNHTCD NLKTCHTLHGSVMAETAVFNR KKLKNSPRIVQSNHLI 21831 52199 B 21959 7 11113							EGSDSRAFAKENNOKAVKETV
QVPQFDQSTIKNIESAKĞLDVW DSWPLQNADGTVAEYNGYHV VFALAGSPKDADTSIYMFVQ KVGDNSIDSWKNAGRVFKDSI KFDANDPILKDTQTQEWSGSAT TSDGKIRLFYTDYSGKHYGKQ LITTAQVNVSKSDDTLKINGVE HKTIFDDRKTYQNGQQFIDER NIPPATYAERPYYFEDKGLKY LGKPTVEPSVGGRLESPWQT, GVSPTVQKLKSLKFDVRQQEE VWEKDEALANWYTPMITCD NLKTCHTLHGSVMAETAVFNP KKLKNSPRIVQSNHLI 21831 52199 B 21959 7 11113		2130	Γ.	1	ľ		
DSWPLONADGTVAEYNGYHV VFALAGSPKDADDTSIYMFYQ KVODNSIDSWKNAGRVFKDSI KFDANDPILKDQTQEWSGSAT TSDGKIRLFYTDYSGKHYGKQ LTTAQVNVSKSDDTILKINGVEI HKTIFDCDGRKTYQNGQQFIDE NIPPATYAERPYFEDKGLKY LGKPTVEPSVCGGRLESPWQT. GVSPTVQKLKSLKFDVRGQEE VWEKDEALANWTYPNHTCD NLKTCHTLHGSVMAETAVFNR KKLKNSPRIVQSNHLI	İ		1				
VPALAGSPKDADDTSIVMFYQ KVGDNSIDSWKNAGRVFKDSI KFDANDPILKDQTQEWSGSAT TSDGKIRLFYTDYSGKHYGKQ LTTAQVNVSKSDDTLKINGVE HKTIPGDGRKTYQNGQFIDE NIPPATPYAERPYYFEDKGLKY LCKPTVEPSVCGGRLESPWQT, GVSPTVQKLKSLKFDVRQQEE VWEKDEALANWTVPNHTCD NLKTCHTLHGSVMAETAVFNP KKLKNSPRIVQSNHLI 21831 52199 B 21959 7 11113	1		l				
KVODNSIDSWKNAGRVFKDSE KFDANDPILKDQTQEWSGSAT TSDGKRLFYTDYSGKHYGKQ LTTAQVNYSKSDDTLKINGVE! HKTIFDGDRKTYQNGQFIDE NIPPATPYAERPYYFEDKGLKY LGKPTVEPSVCGGRLESPWQT. GVSPTVQKLKSLKFDVRGQEE VWEKDEAALANWTVPNHTCD NLKTCHTLHGSVMAETAVFNR KKLKNSPRIVQSNHLI 21831 52199 B 21959 7 11113	1						
KFDANDPILKDQTOEWSGSAT TSDGKIRLFYTDYSGKHYGKQ LITTAQVNVSKSDDTLKINGVEI HKTIFDGDRKTYQNGQGEIDE NIPPATPYAERPYFEDKGLKY LGKPTVEPSVCGGRLESPWQT. GVSPTVQKLKSLKFDVRGQEE VWEKDEALANWTYPNITCD NLKTCHTLHGSVMAETAVFNR KKLKNSPRIVQSNHLI 21831 52199 B 21959 7 11113	ł	l					
TSDGKIRLFYTDYSGKHYGKQ LTTAQVNVSKSDDTLKINGVEI HKTIFODDRKTYQNQQFIDE NIPPATPYAERPYYFEDKOLKY LGKPTVEPSVGGRLESPWOT, GVSPTVQKLKSLKFDVRQQEE VWEKDEALANWYTVPHITCD NLKTCHTLHGSVMAETAVFNP KKLKNSPRIVQSNHLI 21831 52199 B 21959 7 11113						l .	
LTTAQVNVSKSDDTLKINGVĒI HKTIFŪGDKRTV QNGQĢĪDEK NIPPATPYAERPYYFEDKOLKY LGKPTVEPSVCGGRLESPWQT, GVSPTVQKLKSLKFDVSKQĢEE VWĒKDĒAALANWTVPNHTCD NLKTCHTLHGSVMAĒTAVFNR KKLKNSPRIVQSNHLI 21831 52199 B 21959 7 11113	1	1				Ĭ	
HKTIFDODRKTYQNGQQFIDE NIPPATPYAERPYYFEDKOLKY LCKPTVEPSVCGGRLESPWQT. GVSPTVQKLKSLKFDVRQQEE VWEKDEALANWYTPMITCD NLKTCHTLHGSVMAETAVFNP KKLKNSPRIVQSNHLI 21831 52199 B 21959 7 11113	1						
NIPPATPYAERPYYFEDKGLKY LCKPTVEPSVCGGRLESPWQT. GVSPTVQKLSKLKFDVKQGE VWEKDEAALANWTVPNHTCD NLKTCHTLHGSVMAETAVFNP KKLKNSPRIVQSNHLI 21831 52199 B 21959 7 1113						ł	
LGKPTVEPSVCGGRLESPWQT, GVSPTVQKLKSLKFDVRGQEE VWEKDEALANWYTVPNHTCO NLKTCHTLHGSVMAETAVFNP KKLKNSPRIVQSNHLI 21831 52199 B 21959 7 1113	1	1	1	l			
GVSPTVQKLKSLKFDVRGQEE VWEKDEAALANWTVPNHTCD NLKTCHTLHGSVMAETAVFNP KKLKNSPRIVQSNHLI 21831 52199 B 21959 7 11113				ĺ			
VWEKDEAALANWTVPNHTCD NLKTCHTLHGSVMAETAVFNP KKLKNSPRIVQSNHLI		1		l	1		
NLKTCHTLHGSVMAETAVFNP KKLKNSPRIVQSNHLI 21831 32199 B 21959 7				l			
KKLKNSPRIVQSNHLI 21831 52199 B 21959 7 1113	l			l	[
21831 52199 B 21959 7 III3				1			NLKTCHTLHGSVMAETAVFNP
			L				KKLKNSPR\IVQSNHLI
21832 52200 B 21960 1 1188							
	21832	52200	В	21960	1	1188	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21833	52201	В	21961	1	1414	
21834	52202	В	21962	1	1515	
21835	52203	Α	21963	1652	1916	RKNPFILH*LFR*TLRQTKPDNS AGKCVKI**HTQNQRSGRSQND F*RR
21836	52204	Α	21964	1	1353	
21837	52205	A	21965		1507	MAAQDAMYTRISSQ TIDCKEPC LKWAGVVLMLLCWLMFQVL WEACQOKTGEHKTMLQMILCN KSYQQLRLVITPMYLMLGDWL SKNCGQNSTPSLGDNPPERQTO EWSGSATFTSDGKIRLFYTDVS GKHYGKQSLTTAQVNVSKSDD TLKINGYVEDHKTIFDGDGKTYC NVQQFIDEGNYTSGDNHTLLDI HYVEDKGHKYLVFEANTGT GYQGESLFNKAYYGGGTNFF KKESÇKLQSAKKRDAELANG ALGIIELNNDYTLKEVMFLITS NTVTDEIERANVFKMNGKWYL TDSRGSKMTIDDLHHAEDTL LPFYLGEKDDVTYAIKPTCWPC LDIIPSCLALHRIETELMGKFDE GKLPTDPHLMLRJAIETVAHPP AEK*HQNHSSPF*PRFFAVSLRI GSSYHCLISS
21838	52206	A	21966	729	843	NKRIFPSDVKVAEPDHSCV*SFF IGSLASNLSLLINM
21839	52207	В	21967	1	1416	
21840	52208	Α	21968	1	1794	
21841	52209	В	21969	1	1171	
21842	52210	A	21970	1108	1250	NKRIFPSDVKVAEPDHSCV*SFI IGSLASNLSLSLKTRPGGASRVF K
21843	52211	Α	21971	1	2535	
21844	52212	A	21972	743	898	NKRIFPSDVKVAEPDHSCV*SFF IGSLASNLSLSLKTRPAFFQLSIE LSAT
21845	52213	В	21973	1	1605	
21846	52214	В	21974	1	1873	
21847	52215	В	21975	1	2946	
21848	52216	Α	21976	1419	1498	
21849	52217	Α	21977	1	2314	
21850	52218	A	21978	1	1215	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
21851	52219	Α	21979	i	1851	PWISAPVPVDVVEGAMDSVTV
						LSFGGLMLYFCAGWPPARRWC
						FPESISCGSMERDQWWGLQVA
						KRAGLAGGOSGRTVLRERVRIE
						IASTHIALAARHSDWRCCRNGR
						YPARGPAALQNFQRYTGIQHV
						HRIGMAERMWCDRNRERHTVS
						SSGGNRLPNPGPDRSCDNLKTC
						HTSHGSVMAETAVINHKKRKN
						SPRIVQSNDLTEAAYSLSRDQK
						RMLYLFVDQIRKSDGTLQEHD
						GICEIHVA\KYVEIF\GLPFAEAS
				ŀ		KDIRQALKSFAGKEVVFYRPEE
						DAGDEKGYESFPWFIKRAPSPS
l						KGLYSVHINPYLIPFFIGLQN\RL
Į.						TQFRLSETKEIPNPYAM/RLYTN
1						TLCQYRKPDGSGIVSLKIDWIIE
1						RYQLPQSYQRVWTPGINCPEPC
1						PGVWTPGINCPGA WGIHGPHGS
İ		l				AENTGPLHPGQTSAQLETSLLP
1						SLRSARELELAQLSPRRKQNRS
ŀ						LQNIGVTPSLRVPWPKASNVQQ
ŀ		1	1			FIDEGNYTSGDNHTLRDPHYVE
ļ		1	ŀ			DKGHKYLVFEANTGTENGYQG
		1				EESLFNKAYYGGGTNFFRKESQ
ŀ	ĺ	١.				KLQQSAKKRDAELANGALGIIE
		ł				LNNDYTLKKVMKPLITSNTVTD
						EIERANVFKMNGKWYLFTDSR
21852	52220	Α	21980	3197	5376	
21853	52221	Α	21981	543	5690	RKNPFILH*LFR*TLRQTKPDNS
		İ				AGKCVKI**HTQNQRSGRSQND
						F*RRWSRSY
21854	52222	Α	21982	2241	2614	ETLTTLKTKAINTLYSKPTREQ
		1				KTDTKAKNLYLTKRTTAAART
		l				SSVKKAISFNRALKNAMLS*RT
1		l				APSVS*IRHRV*NLTMRSSSSSA
				1		PSPWML*A*AWLCRYCRASCGI
						SSIPTASPVTMACC
21855	52223	Α	21983	31	892	
21856	52224	A	21984	548	855	
21857	52225	В	21985	1	1825	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
01050	52226	_	21986	1	3146	NEW AVENUE NO AVAILABLE
21858	52226	A	21986	1	3146	MEWGKFKRLVLPSQGKSKPEQ
						RCQYRHTVGDQATLPLRSRVE
						KRNGEAEGRIYRKLWDVALLC
						CVQEVINGERKPLRALTEVKSG
		i	İ	İ		FRLREDSDKFDANDPILKDQTQ
						EWSGSATFTSDGKIRLFYTDYS
						GKHYGKQSLTTAQNVQQFIDE
	ŀ					GNYTSGDNHTLRDPHYVEDKG
						HKYLVFEANTGTENGYQGEES
						LFNKAYYGGGTNFFRKESQKL
		1	i			QQSAKKRDAELANGALGIIELN
		<u> </u>				NDYTLKKVMKPLITSNTCDN
21859	52227	A	21987	1	3107	MNEMKREEKFREKRRERNEQI
						LQEIRDYVKRSNLHLIGVSKSD
						GENGTELENTLQDIIQENFPNLA
						RQANIQIQEIQRMLQRYSSRRTT
1		1				PRHIIVRFTKVEMKEKMLRAAR
		l				EKGRVTHKGKPIRLTADLSAET
						LQATKEWRPIFNILKEKNFQPRI
						SYPAKLSFISEGETKYFTDKQM
						LRDFVTTRPALKELLKEALNME
						RNNRLVKSLAWASQNGFRSPA
						LLPLSLMLCAPLQEADGTNLG
						WLAHCGLLCTGDH
21860	52228	В	21988	239	2825	
21861	52229	A	21989	7533	11119	
21862	52230	A	21990	1	3718	MEVIPEGYRLSELKKCSKWIVV
						MVARLYEYTKNQQIIHLKWVN
						HMLGCLLDRNDPLRDVDEAAV
l						PVLCICSADDPVCGPPDHTLTTE
				ŀ	1	LFHSNPYFFLLLSRHGGHCGFL
				İ		RQEPLPAWSHEVILESFRALTEF
1			İ			FRTEERIKGLSRHRASFLGGRRR
ł						GGALQRREVSSSSNLEEIFNWK
l						RSYTRLMAAAAGAAAAPGSRE
				l		PQDRPECGAGHPGPRYYRHPER
						WLLRPEAFLGPLRTRAPSAEDS
						QRERPAARSGPEM
21863	52231	A	21991	1	697	
21864	52232	A	21992	1	978	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
21865	52233	A	21993	145	1929	VSCVUEIADGSERRKA AALTES DYRVI.VGELDDEQMAAL SRLG NDYRVI.SGLDDEQMAAL SRLG NDYRTISA YERGGRYASKLQN EFAGRISALADAECDNILKTCHT SHOSWMAETAVINHKKRKNSP RIVQSNDLTEAAYSI.SRDQKRS RIVQSNDLTEAAYSI.SRDQKRS LYLFVDQIRKSQGTLQEHQBIC EHVAKYAEIFGLTSAEASKDIR QALKSFAGKEVVFYREEDAG DEGGYESFPWIK/RPSRGLYSV HINPYLIPFFIGLQNRFTOFELS ETKEITRPYAMRLYESLCQYRK PDGSGIVSLKIDWIIERYQLPQS YQRMPDFRRRFLQGFCRTRNII RQTGSPAGANQRGPLAATI.SG PGGEGQSAVARLTICEKKNIPIG ETWAGAGSRAALATICHKINIPIG ETWAGAGSRAALATICHKINIPIG ETWAGSERNAINDDVTYAIK PTCWFGLDIIPSCLAHRIETEL TVAHDYDVIVIDSAPNLGIGTIN TVCAADVLHYFTAELEDYTSA LQFFDMLRDLLKNVDLKGNSN GSQSPWMEEQIRD AWGSMYLK NVYRETDEVGKGQIRGNRTYFE QAIDQRSSTGAWRNALSIWERE CNEISIGVSLEDGGSSVELEK

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stup codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	Sequence		031510,211	sequence	or prepriet orquities	
		_	101001	l v v o v	lana	Decorate Market Control
21866	52234	A	21994	I184	2229	RSSVATVALLHAVINHK/KRKN
		1				SPRIVQSNDLTEAAYSLSRDQK
		1				RMLYLFVDQIRKSDGTLQEHD
1		1				GICEIHVAKYAEIFGLTSAEASK
		1				DIRQALKSFAGKEVVFYRPEED
1		1				AGDEKGYESFPWFIKRAHSPSR
1						GLYSVHINPYLIPFFIGLQNRFT
		ł				QFRLSETKEITNPYAMRLYESL
		l				CQYRKPDGSGIVSLKIDWIIERY
		1				QLPQSYQRMPDFRRRFLQVCV
		l				NEINSRTPMRLSYIEKKKGRQT
		l				THIFFDMLRDLLKNVDLKGFEP
		l				DVRILLTKYSNSNGSQSPWMEE
						QIRDAWGSMVLKNVVRETDEV
i		1				GKDTSLSTPAAPMVDSLIARVG
		1				VMARGNAITLPVCGRDVKFTL
		1				EVLRGDSVEKTSRVWSGAVSQ
1		1				GGESFIYMPLTGAAAFLSEMPC
		l				PERRNLKKQCGYSGFAACGYQ
						SQERKNSPRIVQSNDLTEAAYS
		1				LSRDQKRMLYLFVDQIRKSDGT
		l				LQEHDGICEIHVAKYAEIFGLTS
		1		i		AEASKDIRQALKSFAGKEVVFY
			1			RPEEDAGDEKGYESFPWFIKRA
		1				HSPSRGLYSVHINPYLIPFFIGLQ
						NRFTQFRLSETKEITNPYAMRL
		ļ.		ŀ		YESLCQYRKPDGSGIVSLKIDWI
						IERYQLPQSYQRMPDFRRRFLQ
				1		DHNSSPEREQNWMENEFDELT
-				1		EVGFRRWVITNSSELKEHVLTQ
						CKEAKNLDKRLQELLTRISNLE
						NIDDLMELKNTAQELHETYTSI
21867	52235	Α	21995	82	325	
21868	52236	В	21996	220	1173	
21869	52237	A	21997	I	1599	
21870	52238	A	21998	188	421	LO MEN ACTOVICE OF THE PROPERTY OF THE PROPERT
21871	52239	A	21999]1	596	MGMTDMGKSKKKSQGPPSFPP
		1				GGAPPPAIGAPAGRERPPGTGD
		1				PFNQRPCDNLKTCHTSHGSVM
						AETAVINHKKRKNSPRIVQSND
		1		1		LTEAAYSLSRDQKRMLYLFVD
				İ	1	QIRKSDGTLQEHDGICEIHVAK
	I	1				YAEIFGLTSAEASKDIRQALKSF
		1		1		AGKEVVFYRPGEDAGHEKGYE
1		1		1		SFPW/FIKRSPE*RRGKFPFTQM
L						GQD
21872	52240	A	22000	I	1527	
21873	52241	Α	22001	2	870	

SEQ ID		Met	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	of peptide sequence	nod	in USSN 09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21874	52242	Α	22002	I	4304	MAETAVINHKKRKNSPRIVQSN
						DLTEAAYSLSRDQKRMLYLFV
						DQIRKSDGTLQEHDGICEIHVA
			į		İ	KYAEIFGLTSAEASKDIRQALKS
					1	FAGKEVVFYRPEEDAGDEKGY
					İ	ESFPWFIKRAHSPSRGLYSNRFT
					1	QFRLSETKEITNPYAMRLYESL
					İ	CQYRKPDGSGIVSLKIDWIIERY
		1			l	QLPQSYQRMPDFRRRFLQCINA
		ĺ				GHEMTKAIAIAQFNDDSPEARK
		1				ITRRWRIGEAADLVGVSSQAIR
		L				DAEKAGRLPHPD
21875	52243	Α	22003	740	1159	SLPNLDNSAICSSSSSPTRTR*SL
						SVSAAALRRRLPSAISMTPDTL
		1				RPNAGVC*PVSRKEGMRSSSAS
		1				SVSSSWSRSLPDHTREVFSTLSP
		1				RSTSRVNFTSRPHTGKVMALPR
		1				AITPTRAINESTIETAGEDNEVSS
						TG
21876	52244	A	22004	664	1844	SGNLKACHPSHGSVMAETA\VI
						NHKKRKNSPR\IEQ*KDGYQQA
				l		YSLSRDQKRMLYLFVDQIRKSD
						GTLQEHD\GI\CQIHGAQYR*QIS
1						D*PFAEASKDIRQALKSFAGKE
				1		VVFYRPEEDAGDEKGYESFPW
				1		FIKRAHSPSRGLFSGHIPP\YLIPF
						FIGVKNRLTQI\RLFETKEIPHPS
1						AL\RYYESLGQDRK\PDGSSIGS
1		İ				LK\IDWIIERYQLPQSYQRMPDF
						RRRFLQHFLAHAGCPKGTQNFT FRMKFIIFSIKKVLPPVDTILAHD
						PVIHTCHLRESFGSVSPSSPKSN
	ŀ	ļ				RLPCPEVILFHYGESWNLLRAD
1	ŀ					ORLIFAKSWPRASRYQQGHQD
						LFILRSDLPSQVFIRDKLMERRN
						RRTGRTEKARIWGSDGQNGQE
						PGYVEAGCPPLLLTV
21877	52245	A	22005	3	272	POTVEAGETTEELTV
21877	52245	A	22006	I	528	
21879	52247	A	22007	461	831	LLNLATWRYLGTLTRGTAS\VA
210/9	32247	ľ^	22007	101	051	VAGILAALRITNNKLSNHVFVF
				1	1	OGAGEAAMGIAHLLVMALRRK
1				1		K\GTERTTATRKIWMVDSK\GLI
				1		COGEGSHLN/HHEKEMFAODH
	1			1		PEVNSLEEVVRLVKPTAM
	L			L	L	I LYMODELY VILLY KTIAM

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21880	52248	Α	22008	3	1894	EGEDRGLPRTMGAALGTGTRL
		1			1	APWPGRACGALPRWTPTAPAO
	ļ		ĺ			GCHSKPGPARPVPLKKRGYDV
		l				TRNPHLNKGMAFTLEERLOLGI
	İ	ļ				HGLIPPCFLSQDVQLLRIMRYY
						ERQQSDLDKYIILMTLQDRNEK
		1			1	LFYRVLTSDVEKFMPIVYTPTV
		l				GLACQHYGLTFRRPRGLFITIHE
		l				KGHLATMLNSWPEDNIKAVVV
		İ				TDGERILGLGDLGCYGMGIPVG
-		l				KLALYTACGGVNPQQCLPVLL
		l				DVGTNNEELLRDPL\YIGLKHO
		l				RVHGKAYDDLLDEFMQAVTD
		l				KFGINC\LI\QFEDFANANAFRLL
		l				NKYRNKYCMFNDDIQGTASVA
		l				VAGILAALRITNNKLSNHVFVF
		l				OGAGEAAMG\IAHLLVMALE\K
l		l				EGVPKA\EATRKIW\MVDF\KGL
						IVKGRSHLNHEKEMFAQD\HPE
		l				SNSLDEVVRLVKPTAIIGVAAIA
						EA\FTEQILRNMASF\RRAPIIFA
		1				LSNPPRKAECTA\EKCYRVTEG
		l				PRGFFASGKSFLRSVDL/ESMGK
		l				TFIPGGRGNNA*RVPRGWOLG
		l				VHSPGGDPGHIP\DEIFLPDSRA
		l				KLPQEVSEQHL\SQGRLYP\PLS
		l		1		TIRDVSLRIAIKVLDYAYKHNL
						V\SYYPEPKDKEAFCKIPGSYTP
						DYDSFYT/VDSYIWAQGKAMN
21881	52249	Α	22009	322	567	
21882	52250	C	22010	338	603	
21883	52251	A	22011	127	417	
21884	52252	A	22012	253 124	424 1235	M.C. I. DV.C. C. C. D.D. A.I. C. D.T.C. C. W.D.
21885	52253	A	22013	124	1233	YLCLLPKGQGDRALSRTSQWP
		ł				QMSQSQACGGSEQIPGIDIQLN
		1		i		RKYHTTRKLSTTKRFPTACWR
		1				KKVGAFTKIIEAMGFTGPLKYS
		l				KWEDLRLRALRIVY*LCGRKLD
		l				FEGFFSK\CS\MPDTFNSWFLINP
		l				YFHVWMCLVRMKQEGRSGKY
l		l				MCRIIVHFMWEDVQQRGRVM
		l	İ			GVNPYILKKNMILMTNHFYA\A
		l				ILGYDEGILSDDHGLAAALWK
		1		1		NLL\N\RKCEDPRHLELLVEYVR
		1		1		KQIQYLDS\MNGEDLLLTGEVS
	1		1			WRPLVEKNPQSILKPHSPTYND
		l				RGTFDGLGPPHGPPAGFRGTSR
ĺ				1		EEKCLVGPRTPARKWPELDLLN
						KHLSKYLAPICVEFPLSVPQESD
						LLGYRAWENPLALPGCPSSLRG

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21886	52254	A	22014	282	435	GGGASPAACSGSPDSS\HSATA GTGCRRPARPPPAPPTPTPWRS ROOGSRAS
21887	52255	A	22015	3	399	IVALPFGHHHSQTSPLPNLLQTA ACLLLHFLLLSGVETGRPGPWQ PRIRARGPPAPPGPARPAKPRRR /GCRGPAREACWYGCRRPARPP PAPPTPTHGAAGSREAERASEL/ C*GARPLRLGSAGTIMVSACVP
21888	52256	A	22016	279	872	KIPIISLSKLIQGASSKIIPTPQNF HLSPGTKRDSSFQPWSLSAFIIT GQCCSYIVRWGLWSPRULLVV KKYAALSAEMLLVKYEEPSPA ACSGSPDSSSFCNCW*PPAAGR L/ASAAQRPPTPNAHKRSRQQG SRASERAS/CKGRGRYGLGALG GRGGRALGGSRWPPPLPGETLF SGCKHRRRRGSDAAPGEAG
21889	52257	А	22017	106	398	NAAPFSRHAAFQGFSWAVCIFR LPAERALRLFPASCLVLPGGPS GVAPHGWPLPTTAPPLLQGVW RVEVGEHEMAEIFSSAKKEAAG ALQRPCIPVW

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \-possible nucleotide insertion)
21890	52258	Α	22018	I	1806	MRGNTAGGPGDEKPDSTHPPW
						EGTLQEAQEMKKVVRFMWTS
						GGPDPOHWRGSGLYGOLVAO
l						MLHIDVAOVYMDSWWPRCST
l						LAWLRFIWTAGGPDAPHWLRS
						GLGGOLVAWMLHVGVAOVYV
l						DSWWPSCSASMWLSFMEHKO
						CLTLGYEVALFLQHFLSRAFVL
						VWVPRKEKAQVLROGLSLTFR
						ENPGVCVMSSYRPWLITAAAD
						MQPSKPGGTGSYCSYWSPTDN
						TPRLAAACTWNLGSSHGSAPG
						VRTFIEAGFSPPFLAPKALALTL
		ļ				SDSDSRWGPNPVYAQPQP/PEP
						AWGPNPV*GQPQPRNRPG\PEP
						CVCSTPTAEPPWGPNPVSTPTA
1						EPA WGPNPVYA/PTPTAEPPWG
						PNPV*GQPQP/REPAWGPNPVY
						GQPQPRNRPGS\PPQGNLYCSA
	ŀ					ALGATDPSQSWHQCSQNLLPA
						DKGAKREAPW/PRPPSGEVDHQ
l						VWHLHGWPLPTTAPPLLQGVW
						RVEQTPASHHRQPRPRQSSPST
1						KAPPVITVNQGPASHHRQPRPR
l						QSSPSTKAPPVITVNQGPASHH
		İ	1			RQPRPRQSSPST/TGPASHHRQP
						RPRQSSPSTKAPPVITVNQGPAS
						HHRQPRPRQSSPSTKAPPVITVN
	1		1	1		QGPASHHRQPRPRQSSPSTKAP
						PVITVNQGPASHHRQPRPRQSSP
l	l	l				STKAPPVITVNQGPASHHRQPR
l	1					PRQSSTLTKAPPVITVNQSPASH

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21891	52259	A	22019	27	1367	RILAGPARAEAGSASDRSTGHL
1000	02237	1			1.507	FPRGVGMDLCHPQPAELSSGET
		1				DELPRI*WHRKOLLEDIOKLKD
		1				EIADVFAQIDCFESAGDSRMAH
						KENELCIGRKKFNMDPROGRW
		l				LWRARATGCGGCMGAPAW\FL
	l			1		SGEGLNKTAIGTYLGER*ERVE
		1	ľ			P*RGRK*RCAGVYRCVDVQAL
		l				R*VVLGQGPGPQGWHSPSGWG
						CRDPRYCLCNPGVFQSTGARRG
		1				VGPRAPGPLQHCQA*ISQTRPPF
		l				ERFVSMNRGINNGSDLPEDQLR
		1				VREAQPTQPGPGESRRDLSLDG
1		l				WPFFNPDREGWLLKLGERPTD
	İ					TRRRTHTHARAGTHRCMHTTV
1						SRVLLGLPLPPPLPAQPTSQ/PPT
		1				SRC/LRLQDPE/CWSHWEIS*TE
						RGESRVVEGKHESYRISATSAE
ĺ				1		ERDQWIESIR*GVPRSGDGFPSP
						SRRPCFFWVLRDSWRWHWGL
						VTLRVPSPAAPGDP
21892	52260	Α	22020	140	356	
21893	52261	A	22021	2	1147	APARGPAVASGPSPGPAAEPVQ
		l				NTETAAMTTHVTLEDALSNVD
		l				LLEELPLPDQQPCIEPPPSSIMYQ
		l				ANFDTNFEDRNAFVTGIARYIE
		1				QATVHSSMNEMLEEGHEYAV
		1				MLYTWRSCSRAIPQVKCNEQP
		1				NRVEIYEKTVEVLEPEVTKLMK
						FMYFQRKAIERFCSEVKRLCHA
	1	l				ERRKDFVSEAYLLTLGKFINMF
		l				AVLDELKNMKCSVKNDHSAY
	1	1		l		KRAAQFLRKMADPQSIQESQNL
						SMFLANHNRITQCLHQQLEVIP
	1	1		I		GYEELLADIVNICVDYYENKM
1	1	1		I	1	YLTPSEKHMLLKVMGFGLYLM
	1	1				DGNVSNIYKLDAKKRINLSKID
						KFFK/HAAGGAPFRRHADRAGQ
	1	1		1	I	IH*DQCSL*REQVQVDVHPEQH
1	1	ı	l			QPPVQYLRADGSDPG

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence	1	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \-possible nuclcotide insertion)
		1				
21894	52262	Α	22022	47	1581	ALVLMYLASSICMSPKRGTTCS
1		1		1		CLERGSRCPTSEPPPPPEPPPGRS
1		1		i		SRCRPGLGPYGGGGEKMADDP
						SAADRNVEIWKIKKLIKSLEAA
ŀ						RGNGTSMISLIIPPKDQISRVAK
		1		1		MLADEFGTASNIKSRVNRLSVL
		1		1		GAITSVQQRLKLYNKVPPNGLV
		1				VYCGTIVTEEGKEKKVNIDFEP
		1				FKPINTSLYLCDNKFHTEALTA
		1				LLSDDRQVWDFIVIGG/SVGALF
		1		1		GHTPKGNTRGGPWHKFTVGIFP
		1				K\KHG*EGGQ\SALRFARLRME
1		1				KRHNYVRKVAETAVQLFISGD
1		1	1			KVNVAGLVLAGSADFKTELSQ
						SDMFDQRLQSKVLKLVDISYG
				1		GENGFNQAIELSTEVLSNVKFIQ
		ĺ		1		EKKLIGRYFDEISQ/DLTGKYCF
ļ		1		1		GVEDTLKALEMGAVEN\LIVYE
1				l		NL\DIMRYVLHCQGT\KEEKILY
		1		1		LTPEQEK\DKSPFH\DKETGQE\L
		1				ELIESMPLLEWFANNYKKFGA\
	1	1		1	i	TLEMFTDKSQKGSPV**KDLGG
		1				IGG\ILRYQSRFPGEWNTKKET
				l		MNFLTLMTTR
21895	52263	Α	22023	1	296	VEDERVCAGPA WERQAGQERP
						ARKVQGKAPEAEGKHGCLLDI
	i	1		1		LVCTYLAEKGLSGSCGLTEALV
1		1				AFGEGSSLGRKNCRDPWGSLR
						QEPHWSDP/WSGPPERCKGKPQ
1						RRRGSTDAFWTSWSART*PRRG
		1		1		SLAAVG*QRHWWPLGRAAV*E
						ERTAETPGAPSGRSLTGQTLGS
						LHV
21896	52264	A	22024	5	309	LPDADLFCPACGAAAGRARSA
						GMRPVFAVSVARTPSPR/PPPAE
						PQFPRLPCGSIGSPFFRDIDTIQK
						FPDMLSSFFFRDIDTIQKFPDML
						VFNCCCLLNQSR
21897	52265	A	22025	1	258	RGRSRIARSASGPRPALVRPPAR
						PPAP*TPRSWSCWSSC*PRSASA
				1		TGSPSA*ATDAHADSSKAMLPE
		1		1		PTSSISKFSTLQTVPFRL
21898	52266	Α	22026	119	692	PSSHSVRTATYSSLRQSAVSTRT
	1	1				STTTTLAFMARAGGRAGGRAR
1		1	l	1		VGGRTPSGQCGCPACGAAAGR
		1	l	1	1	ARSAGMRPVFAVSVARTPSPR/
	1	1	l	1	1	PPPAEPQFPRRPQR\GAPAPSGG
	1	i	l	1	İ	GASLPAPRALASSIPAQLMKPG
						RAKGSAGNLHPG\EGRPGARRS
		1				DPAGAARSLACP*IWPKQTPSP
	1	1	l	1		LASSSSFFLSETPECAH

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \possible nucleotide insertion)
21899	52267	A	22027	2	246	
21900	52268	Α	22028	1	927	
21901	52269	A	22029	251	1077	VTMGKVLAVSSLPAGPPPPPVP
		l				ALVGLPPPPP*HPGFRLPPLGGG
		1				LGARTSVGRGLERTPGAATASA
		1				AGGAEDGACRCLPNPCTFEECH
		1		ŀ		RKCKEL*MEGVKFTENKAL/SS
		1				KMAIQTQQSKFVKWQVDGEYR
ĺ		1				GSDFTVDVTLWNPDVLMGSGI
		ļ		l		LVAHYLKSITPCLALGSELVYH
		ł				RRPGEEGTVMSVAGKYTLNNW
		l				LATVDVEFEASTRMQDTNVSF
		i				GYQLDLPKANLLFKGSVDSNW
		i				IVGATLEKKLPPLPLTLALGAFL
		l				NHRKNKLRSGFGLTIS
21902	52270	A	22030	1	1171	GGGSGFGCAWRTGWERSPGRE
		1				QAPPPATMGNVLAASSPPAGPP
					1	PPPAPALVGLPPPPPSPPGFTLPP
		l				LGGSLGAGTSTSRSSERTPGAA
						TASASGAAEDGACGCLPNPGTF
		1				EECHRKCKELFPIQMEGVKLTV
				1		NKGLSNHFQVNHTVALSTIGES
		l				NYHFGVTYVGTKQLSPTEAFPV
						LVGDMDNSGSLNAQVIHQLGP
						GLRSKMAIQTQQSKFVNWQVD
						GEYRGSDFTAAVTLGNPDVLV
				1		GSGILVAHYLQSITPCLALGGEL
						VYHRRPGEEGTVMSLAGKYTL
						NNWLATVTLGQAGMHATYYH
						KASDQLQVGVEFE\ASTRMQDT
	ľ					SVSFGYQLDLPKANLLFKGSVD
ł						SNWIVGATLEKKLPPLPLTLAL
						GAFLNHRKNKFQCGFGLTIG
21903	52271	Α	22031	14	434	CYCQGILSPPPCPPFLLGSAQTS
					1	AICSWVRGVPAESSTSSFPKAP*
				1	1	VALESRGEAPTLHLPVTSRFLQ
		1		1	1	CVSSEQECVPWKP/HSSPRPRFL
		1			1	DISPSPTCVRL*APGPVPFDPHQ
1		1			1	KTWRSPLGTRAQGVGPRDRDS
i	1	1		1	1	CRIVPG

SEQ ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21904	52272	Α	22032	989	1859	VYGDSRWSLALSPRLECSGA\IS
						AH*NLCLPGSSDSPASRITGISG
						ARHHAQLI/L/YVFLVEMRFHH
						VGQAGLKLLTSGNP/HHLGLPK
						CWDYSRKPPPRPPRPII\FSFFFL
						RWSLALVAQAGMQWDDLSSL
						QPLPPGFK*FSCLSLLRSWDYG
	ļ					CPPPRLANFVCVCVCVLV/VDG
						GFTMLARLGLELLTSGDLPPFG
				1		LPKCLGFTGMSHCARPIFFF\FE
						MESCFATQAGVQWCDLSSLQP
						LPPRFK*FSCLSLLSSWDYRHM
l						PPCLANF/CFGIFSRDRVSSCWP
21005	60077	١.	22022	157	376	GRSQTPDLK
21905	52273	A	22033	157	370	SHQDFMRTAANLLQTGWCTGS NTGAGARWGLCRASLSRCRCA
						APSAPCSGSRCS*RSWKGTHPS
						LPPRCRGRS
21906	52274	A	22034	298	399	LITRERGRS
21907	52275	À	22035	1	185	
21908	52276	A	22036	311	516	WKAERRTRHLLQTGWCTGSNT
		ľ.				GAGARWGLCRASLSRCRCAAP
	ŀ					SAPRSGSRCS*RLWKGTHPSLPP
l						RCR
21909	52277	Α	22037	I	1487	MMGHSSAIPLTATPGELKGQSP
1						TKMPDPELGCQGAKSQGCSRN
						ARHQKARSMPLQDQHLALAIL
1						LELAVQRGTLSQMLSAILLLLQ
1						LWDSRAQETDNERSAQGTSTL
1						LLSLLQTFQSIICSKDTPPSEGN
1	Ì					MHLLSGPLSPSESFLRESFFTVQ
1						NCRNNEEVTLICKADLENHNK
ł						DGGFWIVIDEKVYDIKDFQTQS
1				i		LTGNSILAQFAGENPVVALEAA FEFEVTRESMHAFCVGQYLEVR
l		ŀ				LYALSDAEDGRGHPLMASVIHL
						LWRLQTSQIHYSYNEEKDEDH
1					i	CSSOWAHLPANRDCSHRWALG
l						DHSQAFLQAIADNNIQDHNVKT
1						HQEQGRSYKEVCTPVIERLRFL
1	1	1	l	l		SNELRPAVGNDLSIISEFKLLSSL
1		1	l	l		PRWRRIAQKIIRERRKKRIPKKP
		1	l	l		ESTADEEKIGNEESDLEEACILP
		1	l	l		HSPINVDKRPIAIKSPKTITSENP
						LG/REPGEHPASPLPPDDAQHA
						HPAAQRKQPRPPAQFRHAGPHS
		1		l		DGTAPDWPQL
		_				

SEQ ID	SEQ ID NO:	Mct	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence	ı	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				scquence		
		L				
21910	52278	Α	22038	263	375	AGPRRMRRISAWAVSTPLTDSP
						SPGHQPA/GEVAGAGSF**HPP/P
						PCPSRYCPQMTPPPPPPAPPPAL
1						PPPPPPAPASPPLPPAPSPPPHAP
						PQALPPPPALPPPPASPPLLPPA
				•		LSPPLPAPP/PPSAPPASPPPAPPP
						PPAPPPSPPPAPPPSAPSSPAPLPP
				i		APASPP\SPAPPPPAPPPAPHPPSP
		ŀ				PAPPCPPLPRPRGPSVPQEKVHQ
						V*IPVP\PAGEGGPAPGRIWPQG
1		ŀ	i	ł		WRRAGKPPTWEEALAAFPPWT
1		ļ.				ALPPASA VPAOPPRAH*VAPAL
						*WALHRRSDSDSDSSNLSPRIPG
						TGGORROPR*DS*APACSEVAG
	ĺ	1				AGSF
21911	52279	A	22039	1	2001	1.051
21912	52280	A	22040	1846	7725	PEPFCHRAAHWSNQYCPFQFQ
21912	32260	<u> </u> ^	22040	1040	1123	DTDSPROTLTVLIKDNGEPSLST
						TATLTVSVTEDSPEARAEFPSGS
1		l				
						APREQKKNLTFYLLLSLILASV
						GFVVTAFGVIIFKVYKWKQSRD
i						LYRAPVSSLYRTPGPSLHADAV
					1	RGGLMSPHLYHQVYLTTDSRR
	ŀ	l				SDPLLKKPGAASPLASRQNTLR
						SCDPVFYRQVLGAESAPPGQIR
	1					YPVPEESQEGTFVGNVAQDFLL
					1	DTDSLSARRLQVAGEVNQRHF
						RVDLDSGALLIKNP
21913	52281	Α	22041	108	340	PQPSQVMSFHGDSGSCRRWAT
1					1	AGT*ATSPPCSSSIFLRRMHSGH
						WCSCWPVRGTPCRDFTAQMAG
1						PSRGSKTNRSMW
21914	52282	Α	22042	275	752	EKTHKTCRCGRGCGKRGTL*H
1				l		CWWECNLLQPLWKMVW/KWL
				!		KELKVELLCDLAVPL/LGINPEE
						KKSLYEKDTCT/R/LFMAAQLAI
1	l	1	1	l		AKKWNOPKCPSINKWLOKLW
	l			1		YIYMMEYYSAIKRN\LTAFAAT
1	l	1	1			WMRLETIILSQGDPLESTCSRSQ
1				1		ANROEITSVPKA
21915	52283	A	22043	25	202	DAEAWAMELADVGA*ASSQG
21913	34283	l ^A	22043	23	202	
1	l				l	APQPEEPAASLGVNEKGGAGG
						RSGRRPLGRDRPRPTRLL

IQKLSFGEDNSYVILSGPPQSF EDSFKKCSSEVEAKNKIEELL LLNRVCQDGRKPHTVRLIRF SSEKHVGRESRQCHIPSHVIQ GTGNYDVMPPWDLIMKLF MVNVKMPFRLTLLSMCFCSL AVNTATTRLIDYYLMPTISITS SGKHSFKMKDTHMEDFPKDI TNRDFLPSGRIESTRTRESPLE TNFSKEKDINEFPLCSLPEGV EVFKQLPVDIQEELISGKSREF QGKGSVSCPLHASRGVLSFF KQMQDIPINFRDHLSSSKQVS VSPCEPGTSGFNSSSSSYMSS DYSYYLDNRLKDERISQGFK QGFHFTNSNPAIVSAFHSIPF	SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
	NO:		hod				
21916 S2284 A 22044 I 2290 MEKLGVEPEEGGGDDEED: AWAMELADVGAAASSQGAI PEEPAASLGVNEKGVHDQVI PINASSR VITHVDLDCFYAQV MISNPELKDKPLGVVQKYLV TCNYEARKLGVKKLMNVRE KEKCPQLVLVNOEDLTRYRE SYKVTELLEEFSPVVERLOFI NFVDLTEMVERLQQLQSDI SAVTVSGHYVNNQSINLLLDV IRLLVGSQIAAEMREAMYNQ GLTGGAGVASNKLLAKLVS FKPNQOTVLLPESCQHLHISL HIKEIPGIGYKTAKCLEALGN VRDLQTESPKLEKELGISVA IQKLSFGEDNSPVILSOPPQSF EDSFKKCSSEVEAKNIEELL LLNRVCQDGRKPHTVRLIRE SSEKHYGRESRQCHIPSHVQ GTGNYDVMPPNDILMKLF MYNVKMPFRLTLLSNCFCSL AVNTATTRLIOPYMPNISTI SGKHSFKMKDTHMEDFPKDI TNRDFLPSGRIESTRTRESPL TNRSKENDIPFCLSPEGV EVFKQLPVDIQEELISGKSRE QGKGSVSCPLHASRGVLSFS KQMQDIPINPRDHLSSKQVS VSPCEPGTSGFNSSSSSYMSS DYSYYLDNRLKDERSGQFK QGFHFTNSNPAVSAFHSFP QSEQLFSRNHTTDSHKQTVA		sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
AWAMELADVGAAASSQGAI PEPAASLGVNEKGVHDQVI PNASSLGVNEKGVHDQVI PNASSLGVNEKGVHDQVI PNASSLGVNEKGVHDQVI PNASSLGVNEKGVHDQVI PNASSLGVNEKGVHDQVI MISNPELKDKPLGVQGYTLV TCNVEARKLJGVKKLMNVER KEKCPQLVLVNGEDLTRYRE SYKVTELLEFSPVVERLOFL NFVDLTEMVEKRLQQLQSDI SAVTVSGHYVNNQSINLLDV IRLLVGSQIAAGMEAMYKC GLTGCAGVASNKLLAKLVS FKPNQOTVLLIPESCQHLIHSL HIKEIPGIGYKTAKCLEALON VRDLQTFSPKLEKELGISVA IQKLSSGEDNSPVLSGPPQSS EDSFKKCSSEVEAKNKIEELL LLINEVCQDGKFHTVRLIIRE SSEKHVGESRQCHIPSHVIQ GTGNYDVMPPMVDILMKLF MVNVKMPFRLTLLSMCFCSL AVNTATTRLIDYYLMPTISIT SGKHSFKMKDTHMEDFPKDI TNRFOLPSGRIESTETRESPL TNFSKEXDIPEFLCSIPEGV EVFKQLPVDIQEELSGKSREI QGKGSVSCPLHASRGVLSFFE KQMQDIPINFBFLGS.PGEV EVFKQLPVDIQEELSGKSREI QGKGSVSCPLHASRGVLSFFE KQMQDIPINFBFLISSSSVSSS DYSYYLDNRLKDERISGGRE QGKGTSTSTRESSPLT TNFSKEXDIPEFLGS.PGEV SVSTCEFGTSGFNSSSSSYMSS DYSYYLDNRLKDERISGGRE QGFHTNISNPAVSAFHSIFP QSEQLFSRNHTTDSHKQTVA					sequence		
AWAMELADVGAAASSQGAI PEPAASLGVNEKGVHDQVI PNASSRVIVHVDLDCFYAQV MISNPELKDKPLGVQGYYLV TCNYEARKLJGVKKLIMYNEK KEKCPQLVLVNGEDLTRYRR SYKVTELLEFSPYVERLOFE NFVDLTEMYEKRLQQLQSDI SAVTVSGHYVNNQSNLLDV IRLLVGSQIAAGMEAMYNC GLTGCAGVASNKLLAKLVS FKPMQOTVLLPESCQHLIHSL HIKEIPGIGYKTAKCLEALON VRDLQTFSPKLEKELGISVA IQKLSSGEDNSPVLSGPPQSS EDSFKKCSSEVEAKNKIEELL LLNRVCQDGKPHTVRLIRR SSEKHVGESRQCHIPSHVQ GTGNYDVMPPMVDILMKLF MVNVKMPFRLTLLSMCFCSL AVNTATTRLIDYYLMPTISIT SGKHSKMKDTHMEDFPKDI TNRDFLPSGRIESTETRESPL TNFSKEXDIPSFLCSSPCSI QGKGSVSCPLHASRGVLSFF KQMQDIPNTPGDLISSKGVSV VSCEFGTSGFNSSSSSYMSS DYSYYLDNRLKDERISGRE QGKGSVSCPLHASRGVLSFF KQMQDIPNTPGHLSSSKGVS VSVECFGTSGFNSSSSSYMSS DYSYYLDNRLKDERISGRE QGKGTSVSVLSFF KQMQDIPNTPGLISSKGVS VSVECFGTSGFNSSSSSYMSS DYSYYLDNRLKDERISGRE QGKGTSTRINSPRAVSAFHSIFP QSEQLFSRNHTTDSHKQTVA	21016	52284	ΙΔ	22044	lı .	2200	MENI CAEBEECCCODEEDAE
PEPPASLGVNEKGVHÖVI PNASSRVIVHVDLDCFY AQV MISNPELKBKPLGVQQKYL. TCNYEARKLGVKKLMNYEE KEKCPQ.VLV.NORGD.TRYRE SYKVTELLEFSPVVERLGFE NFYDLTEMYEKRLQQLGSDI SAVTVSGHYVNQSNLLDV IRLLVGSQIAAEMREAMYNQ GLTGCAGVASNKLLAKLVSQ FKPNQQTVLLPESCQHLIHSL HIKEIPGIGYKTAKCLEALGR VRDLQTFSFRLERELGISVA IQKLSFGEDNSPVILSGPPQSF EDSFKCSSEVEAKNIEELL LLNRVCQDGRKPHTVRLIIR SSEKHYGRESRQCHIPSHVQ GTGNYDVMPPVDLMKLFF MYNVKMPFRLTLLSNCFCSL AVNTATTRLIGYYLMPTISTT SGKHSFKMKDTHMEDFFXDL TNRFEKENDFFLCSLFGEV EVFKQLPVDIQEELISGKSREI QGKGSVSCPLHASTGVLSFFS KQMQDIPINFRDHLSSSKQVS VSPCEPGTSGFNSSSSYMSS DYSYYLDNRLKDERISGGRS GQFHFTNSNPAVSAFHSFP QSEQLFSRNHTTDSHKQTVA 21917 \$2285 A 22045 2 141	21910	32204	^	22044	ľ	22,50	
PNASSR VIVHVDLDCFY-ÀÖV MISNPELKDKPLGVQQKYLV TCNYEARKLGVKKLMNVRE KEKCPQLVLVNGEDLTRYR SYKYTELLEFSPVYBELGFI NFVDLTEMVEKRLQQLQSDI SAVTVSGHVYNNQSINLLDV IRLLVGSQIAAEMREAMYNG GLTGCAGVASNKLLAKLVS FKPNQGTVLIPESCQHLIHSL HIKEIPGIGYKTAKCLEALGN VRDLQTFSPKILEKELGISVA IQKLSFGEDNSPVLSGPPQSF EDSFKKCSSEVEAKNKIEELL LLNRVQDGRFHTVRLIRR SSEKHVGRESRQCHIPSHVQ GTGNYDVMPPMVDLMKLFI MVNVKMPFRLTLLSMCFCSL AVNTATTR.IDYYLMPTISIT SGKHSKMKDTHMEDFFKD) TNRDFLPSGRIESTRTRESPL TNFSKEKDNEFPLCS.PEGV EVFKQLPVDIQEELSGKSREI QGKGSVSCPLHASRGVLSFF KQMQDIPNTPRDHLISSSKQVS VSPCEPGTSGFNSSSSSYMSS DVSYYLDNRLKDERISGREK QGKGSVSYCHASRGVLSFF KQMQDIPNTPRDHLISSSKQVS VSPCEPGTSGFNSSSSSYMSS DVSYYLDNRLKDERISGREK QGFHFTNSNPAVSAFHSIFP QSEQLFSRNHTTDSHKQTVA					l		
MISNPELKOKPLGVOQOKYLV TCNYEARKLGVKKLMNYRE KEKCPQLVLVNGEDLTRYRE SYKVTELLEFSPVVERLGFI NFVDLTEMVERLQCLQSDI SAVTVSGHVYNNQSINLLDV IRLLVGSQIAAEMREAMYNQ GLTGCAGVASNKLLAKLVS FKPNQOTVLLPESCQHLHISL HIKEIPGIGYKTAKCLEALOR VRDLQTFSPKLEKELGISVA IQKLSFGEDNSPVILSOPPQSF EDSFKKCSSEVEAKNITEELL LLNRVCQDGRKPHTVRLIRE SSEKHYGRESRQCHIPSHVIQ GTGNYDVMPPMVDILMKLFI MYNVKMPFRLTLLSNCFCSL AVNTATTRLIOTYLLIPS SGKHSFKMKDTHMEDFPKDI TINDFLPSGRIESTRTRESPL TNFSKENDIPFCSLPGEVV EVFKQLPVDIQEELISGKSREI QGKGSVSCPLHASRGVLSFS KQMQDIPINPRDHLSSKQVS VSPCEPGTSGFNSSSSSYMSS DYSSYLDNRLKDERSGQRE QGFHFTNISNPAVSAFHSFP QSEQLFSRNHTTDSHKQTVA					l		
TCNYEARKLGVKKLMNVEC KEKCPQLVLVNGEDLTRYRE SYKYTELLEEFSPVVERLGFT NFVDLTEMVEKRLQQLQSDI SAVTVSGHVYNNQSINLLDV IRLLVGSQIAAEMREAMYNC GLTGCAGVASNKLLAKLVSC FKPNQGTVLLFESCQHLIHSL HIKEIPGIGYKTAKCLEALGN VRDLQTFSFKLIFKELGISVA IQKLSFGEDNSPVILSGPPQSF EDSPKKCSSEVEAKNKIEELL LLNRVCQDGRKPHTVRLIFR SSEKHYGRESRQCHIPSHVQ GTGNYDVMPPMVDLMKLF MYNVKMPFRLTLLSNCFCSL AVNTATTRLIDYYLMPTISIT SGKHSFKMKDTHMEDFFKDI TNRDFLPSGRIESTRTRESPL TNRSKENDIFFLCSLPGVV EVFKQLPVDIQEELSGKSREI QGKGSVSCPLHASRGVLSFF KQMQDIPNPRDHLSSSKQVS VSPCEPGTSGFNSSSSSYMSS DYSYYLDNRLKDERISGGRE KQMGDIFNFRDHLSSSKQVS VSPCEPGTSGFNSSSSYMSS DYSYYLDNRLKDERISGGRE GGFHTNSNPAVSAFHSIFP QSEQLFSRNHTTDSHKQTVA							
KEKCPQLVLVNGEDLTRYRE SYKVTELLEFSPVVERLGETE NFVDLTEMVEKRLQQLQSDI SAVTVSGHVYNNQSINLLDV IRLLVGSQIAAEMREAMYNQ GLTGCAGVASNKLLAKLVSS FKPNQOTVLLPSCGHLHISL HIKEIPGIGYKTAKCLEALGI VRDLGTFSFKLEKELGISVA IQKLSFGEDNSPVLSSPQSF EDSFKKCSSEVEAKNKIEELL LLNRVCQDGRKPHTVRLIRE SSEKHYGRESQCHIPSHVQ GTGNYDVMPPMVDILMKLF MYNVKMPFRLTLLSNCFCSL AVNTATTRLDYYLMPTISTI SGKHSFKMKDTHMEDFFKD TNRSKENDINEFPLCSPFGCV EVFKQLPVDIQEELISGKSREI QGKGSVSCPLAASRGVLSFF KQMQDIPNPRDHLSSSKQVS VSCEGTGGFTSGNSSSSSYMSS DYSSYLDNRLKDERISGRAF QGFHFTNSNPAVSAFHSIFP QSEQLFSRNHTTDSHKQTVA							, ,
SYKVTELLEEFSPVVERLGEL NFVDLTEMVEKRLQQLQSDI SAVTVSGHVYNNQSINLLDV IRLLVGSQIAAEMREAMYNC GLTGGAGVASNKLLAKLVS FKPNQQTVLLPESCQHLIHSL HIKEIPGIGYKTAKCLEALGN VRDLQTESFKLEKELGISVA IQKLSFGEDNSPVILSGPPQSF EDSFKKCSSEVEAKNIEELL LLNRVCQDGRKPHTVRLIRE SSEKHVGRESRQCHIPSHVQ GTGNYDVMPPVDLIMKLF MYNVKMPFRLTLLSNCFCSL AVNTATTRLIOTYLMPTISTIS SGKHSFKMKDTHMEDFPKDI TNRDFLPSGRIESTETRESPL TNFSKEKDINEFPLGSLPGEV EVFKQLPVDIQEELSGKSRE QGKGSVSCPLHASRGVLSFS KQMQDIPINPRDHLSSSKQVS VSPCEPGTSGFNSSSSSYMSS DYSSYLDNRLKDERISGPK QGFHFTNSNPAVSAFHSFP QSEQLFSRNHTTDSHKQTVA					İ		
NFVDLTEMVEKRLQQLQSDI SAVTVSGHVYNNQSINLLDV IRLLVGSQIAAEMREAMYNC GLTGCAGVASNKLLAKLVSC GLTGCAGVASNKLLAKLVSC FKPNQGTVLLPESCQHLIHSL HIKEIPGIGYKTAKCLEALGI VRDLQTFSPKILFKELGISVA IQKLSFGEDNSPVILSGPPQSF EDSFKKCSSEVEAKNKIEELL LLNRVQQDGKPHTVRLIHS SSEKHVGRESRQCHIPSHVQ GTGMYDVMPPMVDLIMKLEI MVNVKMPFRLTLLSMCFCSL AVNTATTRLIDYYLMPTISIT SGKHSFKMKDTHMEDFPKD) TNRDFLPSGRIESTRTRESPL TNFSKEXDIPSFLCSLPGGV EVFKQLPVDIQEELSGKSREI QGKGSVSCPLHASRGVLSFF KQMQDIPINFBPLISSSKQV VSPCEPGTSGFNSSSSSYMSS DVSVYLDNRLKDERISGFK QGFHFTNSNPAVSAFHSIFP QSEQLFSRNHTTDSHKQTVA			l				
SAVTVSGHYVNNQSINLIDV IRLLVGSQIAAEMREAMYNQ GLTGCAGVASNKLLAKLVSC FKPNQOTVLLPESCQHLHISL HIKEIPGIGVKTAKCLEALGI VRDLQTFSPKILEKELGISVA IQKLSRGEDNSPVILSGPPQSF EDSFKCSSEVEAKNIEELL LLNRVCQDGRKPHTVRLIRE SSEKHYGRESRQCHIPSHVIQ GTGNYDVMPPMVDILMKLF MYNVKMPFRLTLLSMCFCSL AVNTATTRLIDYYLMPTISTI SGKHSFKMKDTHMEDFPKDI TINFSKEKDINEFPLCSLPEGVV EVFKQLPVDIQEELISGKSREI QGKGSVSCPLHASRGVLSFF KQMQDIPINPRDHLSSSKQVS VSPCEPGTSGFNSSSSSYMSS DYSSYLDNRLKDERISGREE QGKGSVSCPLHASRGVLSFF KQMQDIPINPRDHLSSSKQVS VSPCEPGTSGFNSSSSSYMSS DYSSYLDNRLKDERISGREE QGFHFTNISNPAUVSAFHSIFP QSEQLFSRNHTTDSHKQTVA							
IRLLVSSQIAAEMREAMYNC GLTGCAGVASNKLLAKLVSC FKPNQGTVLLPESCQHLIHSL HIKEIPGIGYKTAKCLEALOR VRDLQTESPKLIFEKELGISVA IQKLSFGEDNSPVILSGPPQSF EDSFKKCSSEVEAKNKIEELL LLNRVQQDGRFHTVRLIRR SSEKHVGRESRQCHIPSHVQ GTGNYDVMPPMVDLIMKLF MYNVKMPFRLTLLSNCFCSL AVNTATTRLIDYYLMPTISIT: SGKHSFKMKDTHMEDFFKDL TNRDFLPSGRIESTRTRESPL TNRSKENDIFFLCSLPGOV EVFKQLPVDIQEELSGKSREI QGKGSVSCPLHASRGVSFS KQMQDIPINPRDHLSSKQVS VSPCEPGTSGFNSSSSSYMSS DYSYYLDNRLKDERISGGRE KQMGPFTNSNPAVSAFHSFP QSEQLFSRNHTTDSHKQTVA			l				
GLTGGAĞVASNKLLAKLVSĞ FKPNQTVLLPESCQHLIHSL HIKEIPGIGVKTAKCLEALGIB VRDLQTFSPKLLEKELGISVA IQKLSFGEDNSPVILSGPPQSF EDSFKKCSSEVEAKNIKIEELL LLNRVQDGRKPHTVRLIRE SSEKHYGRESKQCHIPSHVQ GTGNYDVMPPMVDILMKLFI MVNVKMFRLTLLSMCFCSL AVNTATTRLIDYYLMPTISTI SGKHSFKMKDTHMEDFPKDI TNRDFLPSGRIESTRTRESPLE TNFSKEKDINEFPLCSLPEGVV EVFKQLPVDIQEELISGKSREI QGKGSVSCPLHASRGVLSFF KQMQDIPINPRDHLSSSK QVS VSPCEPGTSGFISSSSSYMSS DYSYYLDNRLKDERISQFRK QGFHFTNSNPALVSAFHSIFP QSEQLFSRNHTTDSHKQTVA			ĺ		l .		
FKPNQQTVLLPESCQHLHSL HIKEIPGIGYKTAKCLEALGN VRDLQTFSFKILEKELGISVA IQKLSFGEDNSPVILSOPPQSF EDSFKCSSEVEAKNIEELL LLNRVCQDGRKPHTVRLIIR SSEKHVGRESRQCHIPSHVQ GTGNYDVMPPNDILMKLF MYNVKMPFRLTLLSNCFCSL AVNTATTRLDYYLMPTISTI SGKHSFKMKDTHMEDFPKDI TNRDFLPSGRIESTRTRESPL TNRSEKDINEFFLCSIPEGV EVFKQLPVDIQEELSGKSRE QGKGSVSCPLHASRGVLSFF KQMQDIPINPRDHLSSSKQVS VSPCEPGTSGFNSSSSSYMSS DYSYYLDNRLKDERISQFK QGFHFTNSNPAUVSAFHSIFP QSEQLFSRNHTTDSHKQTVA			l		l		
HIKEIRĞIGYKTAKCLEALGI VRDLQTFSPKILEKELĞISVA IQKLSFGEDNSPVILSGPPQSF EDSFKKCSSEVEAKNKIEELL LLNRVQQDGKPHTVRLIIR SSEKHYGRESRQCHIPSHVQ GTĞNYDVMPPMVDLIMKLIF MVNVKMPFRLTLLSMCPCSL AVNTATTRLIDYYLMPTISIT SĞKHSFKMKDTMHEDFPKDL TNRDFLPSGRIESTRTRESPLL TNFSKEXDIPSFLCS.PEGV EVFKQLPVDIQEELSĞKSREI QĞKGSVSCPLIASRQVLSFF KQMQDIPINPBDHLSSSKQVS VSPCEPGTSĞFNSSSSSYMSS DYSYYLDNRLKDERISĞĞRK QĞFHFTNSVPAVSAFHSIFP QSEQLESRNHTTDSHKQTVA							
VRDLQTFSPKLEKELGISVA IQKLSFGEDNSPVLSSPPQSF EDSFKKCSSEVEAKNKIEELL LLNRVCQDGRKPHTVRLIRF SSEKHYGRESGQHISPHVQ GTGNYDVMPPMVDILMKLF MYNVKMPFRLTLLSMCFCSL AVNTATTRLIDYYLMPTISTI SGKHSFKMKDTHMEDFPKDI TNRSHEKDINEFPLCSLPEGV EVFKQLPVDIQEELISGKSREI QGKGSVSCPLAASRGVLSFF KQMQDIPINPRDHLSSSKQVS VSPCEPGTSGFNSSSSSYMSS DYSSYYLDNRLKDERSGQFK QGFHFTNSNPAUVSAFHSIFP QSEQLFSRNHTTDSHKQTVA			1				
IQKLSFGEDNSPVILSOPPOSE EDSFKKCSSEVEAKNKIEELL LLNRVQDGRKPHTVRLIRR SSEKHVGRESRQCHIPSHVQ GTGNYDVMPPMVDILMKLF MYNVKMPFRLTLLSNCFCSI AVNTATTRLIDYYLMPTISIT: SGKHSFKMKDTHMEDFFKDI TNRDFLPSGRIESTRTRESPLL TNFSKEKDINEFFLCS.PEGV EVFKQLPVDIQEELSGKSREI QGKGSVSCPLHASRGVLSFFS KQMODIPINPRDHLSSSKQVS VSPCEPGTSGFNSSSSSYMSS DYSYYLDNRLKDERISGGFK QGFHFTNSNPAVSAFHSIFP QSEQLFSRNHTTDSHKQTVA			l				
EDSFKKCSSEVEAKNKIEĒLI LLNRVCQDGRKPHTVRLIRR SSEKHYGRESKQCHIPSHVIQ GTGNYDVMPPMVDILMKLF MVNVKMPFRLTLLSMCFCSL AVNTATTRLIDYYLMPTISTT SGKHSFKMKDTHMEDFPKDI TNRPFLPSGRIESTKTRESPLL TNFSKEKDINEFPLCSLPEGV EVFKQLPVDIQEĒLISKSKEI QGKGSVSCPLAHSRGVLSFF KQMQDIPINPRDHLSSSKQVS VSPCEPGTSGFNSSSSSVMSS DYSYYLDNRLKDERISQFK QGFHFTNSNPAUVSAFHSFP QSEQLFSRNHTTDSHKQTVA					l		VRDLQTFSPKILEKELGISVAQR
LLNRVCQDGRKPHTVRLIRE SSEKHYGRESRQCHIPSHVIQ GTGNYDVMPPMVDILMKLF MYNVKMPFRLTLLSNCFCSI. AVNTATTRLJOYYLMPTISTI SGKHSFKMKDTHMEDFPKDI TINRDFLPSGRIESTRTRESPLE TINFSKEKDINEFPLGS.PEGVV EVFKQLPVDIQEEILSGKSREI QGKGSVSCPLAASGGVLSFS KQMQDIPINPRDHLSSSKQVS VSPCEPGTSGFISSSSSSYMSSI DYSYYLDNRLKDERISQFSK QGFHFTINSNPALVSAFHSFP QSEQLFSRNHTTDSHKQTVA							IQKLSFGEDNSPVILSGPPQSFSE
SSEKHYĞRESRQCHIPSHVIQ GTĞNYDVMPPMVDILMKLFI MYNYKMPFRLTLISMCFCSL AVNTATTRLIDYYLMPTISIT: SGKHSFKMKDTHMEDFPKDI TINRDFLPSGRIESTRTRESPLI TINFSKEXDINEPFLGS.IPEGVI EVFKQLPVDIQEEILSGKSREI QGKGSVSCPLIASRGVLSFF KQMQDIPINPRDFILISSSKQVS VSPCEPGTSGFNSSSSSYMSS DYSYYLDNIKLBERISGÇRK QGFHFTNISIPAVSAFHSIFP QSEQLESRNHTTDSHKQTVA			ĺ				EDSFKKCSSEVEAKNKIEELLAS
GTGNYDVMPPMVDILMKLE MVNVKMPFRLTLISMCFCSL AVNTATTRLIDYYLMPTISTT SGKHSFKMKDTHMEDFPKDI TNRDFLPSGRIESTRTRESPLL TNFSKEKDINEFPLCSLPEGV EVFKQLPVDIQEELISGKSREI QGKGSVSCPLHASRGVLSFS KQMQDIPINPRDHLSSSKQVS VSPCEPTGSGTSSSSSYMSS DYSYYLDNRLKDERISQFK QGFHFTNSNPAUVSAFHSIPP QSEQLFSRNHTTDSHKQTVA 21917 52285 A 22045 2 141			l				LLNRVCQDGRKPHTVRLIIRRY
MYNVKMPFRLTLLSNCFCSL			l				SSEKHYGRESRQCHIPSHVIQKL
AVNTATTRLIDYYLMPTISIT:			1				GTGNYDVMPPMVDILMKLFRN
SGKHSFKMKDTHMEDFPKD TNDBH_PSGRIESTRESPLE			l				MVNVKMPFRLTLLSMCFCSLK
TNRDFLPSGRIESTRTRESPLE TNTSKEKDINEFPLCSLPEGV' EVFKQLPVDIGEELISGKSBEI QGKGSVSCPLHASRGVLSFFS KQMQDIPINPRDHLSSSKQVS VSPCEPGTSGFINSSSSSYMSS DYSYYLDNRLKDERISQGPK QGFHFTNSNPAIVSAFHSIFPI QSEQLFSRNHTTDSHKQTVA							AVNTATTRLIDYYLMPTISITSR
TNFSKEKDINEFPLCSLPEGV EVFKQLPVDIQEELISGKSREI QGKGSVSCPLHASGGVLSFF KQMQDIPINPRDHLSSSKQVS VSPCEPGTSGFNSSSSSYMSS DYSYYLDNIKLKDERISQFK QGFHFTNISNPAUVSAFHSIFP QSEQLFSRNHTTDSHKQTV.A							SGKHSFKMKDTHMEDFPKDKE
EVFKQLPVDIGEELSGKSREI QGKGSVSCPLHASRGVLSFIS KQMQDIPINPRDIHLSSSKQVS VSPCEPGTSGFISSSSSSYMSSI DVSVYLDNIKLKDERISGPK QGFHFTNISVPALVSAFHSIFP QSEQLFSRNHTTDSHKQTVA							TNRDFLPSGRIESTRTRESPLDT
QGKGSVSCPLHASRGVLSFS KQMQDIPINPRDHLSSSKQVS VSPCEPGTSGFINSSSSSYMSS DYSYYLDNRLKDERISQGPK QGFHFTNSNPAIVSAFHSIFPI QSEQLFSRNHTTDSHKQTVA 21917 52285 A 22045 2 141							TNFSKEKDINEFPLCSLPEGVDQ
KQMQDIPINPRDHLSSKQVS VSPCEPGTSGFNSSSSSYMSS DYSYYLDNRLKDERISQFK QGFHFTNSNPAUVSAFHSIFP QSEQLFSRNHTTDSHKQTV.A 21917 52285 A 22045 2 141							EVFKQLPVDIQEEILSGKSREKF
VSPCEPGTSGFNSSSSSYMSS DYSYYLDNRLKDERISGGPK QGFHFTNSNPA\VSAFHSIFPI QSEQLFSRNHTTDSHKQTV-A 21917 52285 A 22045 2 141							QGKGSVSCPLHASRGVLSFFSK
DYSYYLDNRLKDERISQGPK QGFHFTNISNPAIVSAFHSIFPI QSEQLFSRNHTTDSHKQTVA 21917 52285 A 22045 2 141							KQMQDIPINPRDHLSSSKQVSS
QGFHFTNSNPALVSAFHSIFP QSEQLFSRNHTTDSHKQTVA 21917 52285 A 22045 2 141							VSPCEPGTSGFNSSSSSYMSSQK
QGFHFTNSNPALVSAFHSIFP QSEQLFSRNHTTDSHKQTVA 21917 52285 A 22045 2 141			ı				DYSYYLDNRLKDERISOGPKEP
QSEQLFSRNHTTDSHKQTV\A							QGFHFT\NSNPA\VSAFHS\FPNL
							QSEQLFSRNHTTDSHKQTV\AT
21918 52286 C 22046 278 489	21917	52285	A	22045	2	141	
	21918	52286	С	22046	278	489	
21919 52287 A 22047 130 426	21919	52287	Α	22047	130	426	

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nuclcotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		_				
21920	52288	A	22048	357	1579	QRRPRPFPSQGISMTECFLPPTSS
						PSEHRRVEHGSGLTRTPSSEEIS
						PTKFPGLYRTGEPSPPHDILHEP
l .						S*SNGLMDEKDHGKKKGNFKK
		1				KEKRTEGYAAFQEDSSGDEAE
		1				KLMPRFEEAYGRTTETEKVQEF
ŀ						QRLLKELPEGNYLLISLFVVHM
ŀ		l				DHVITKELEAKMNIQNIYIVVSP
l					-	TVQISNLVLYVFFTHVQELFGN
1						VVLKQVTKTRRWSNVATMPTL
						PETQAGFKEEIRIQELLLNCLHQ
	İ					DLQGGIKDLSEEERLWEEKEIL
						TKQEELLGMEQFLRRQIASEKE
	l		1			EIECLRAETAEIQSHQQQGLSET
		1				EEYSSESESEDEEELQIILEDLQR
		l				QNEELEIKNNHLNQAIHEESEA
						VIEPRMQPWLLQLQPDRAKQQ
						AQEDEEPEWRGGAMQTPKNGI
21921	52289	Α	22049	182	2212	VTEKTSSLPSQGISMTECFLPPT
						SSPSEHRR\VEHG/SAGLTPGPPA
						SEEIQPLPKFPGLY\RTGEPPHPP
						P*HPSHEPSLM*CPDDEKDHGK
		1				KKGKFKKKEKRTEGYAAFQED
1		1		i		SSGDEAESPSKMKR\SKGIHVFK
						KPSFSKK\REKDFKIKEKPKEEK
1		1	ŀ			HKEEKHKEEKHKEKKSKDLT\A
1						ADVVKQWKEKKKKKKPIQEPE
1						VPOIDVPNLKPIFGIPLADAVER
						TMMYDGIRLPAVFRECIDYVEK
						YGMKCEGIYRVSGIKSKVDELK
						AAYDREESTNLEDYEPNTVASL
				i		LKOYLRDLPENLLTKELMPRFE
						EACGRTTETEKVQEFQRLLKEL
						PECNYLLISWLIVHMDHVIAKE
	}			İ		LETKMNIONISIVLSPTVOISNR
				!		VLYVFFTHVQELFGNVVLKQV
						MKPLRWSNMATMPTLPETQAG
						IKEEIRROEFLLNCLHRDLOGGI
	1					KDLSKEERLWEVQRILTALKRK
1						LREAKRQECETKIAQEIASLSKE
	1		l			DVSKEEMNENEEVINILLAGEN
			l			EILTEQEELLAMEQFLRRQIASE
1			l		1	KEEIERLRAEIAEIQSRQQHGRS
1			l			ETEEYSSESESESEDEEELQIILE
1			1			
			1			DLQRQNEELEIKNNHLNQAIHE
1			l			EREAIIELRVQLRLLQMQRAKA
			l			EQQAQEDEEPEWRGGAVQPPR
01000	50000	<u>_</u>	02050	2.4	202	DGVLEPKAAKEQPKAGKEPAK
21922	52290	В	22050	34	322	

	01/0/506/					PC1/US01/08631
SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first		Amino acid sequence (X=Unknown,
NO:	sequence	hod	09/540,217	codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	Sequence			sequence	or popular sequence	and the second s
		<u> </u>				
21923	52291	A	22051	137	1546	QMLGGRPPPRWLQLAKARAL
						RRRPPPSLHLTLPTALSLKS/RT
		1				VPKSIPISETPNIPPVSVQPPASI
						PPLGVPP/PEPSHGDDQPRPG\F
						PIFMEQQIMQQIRPPFIRGPP\PI
						PPTPTAPCPTPCFPASGPRPVA
						ETWAPLPAPCTGPCY\PPHPPP
						PPPP/SPGNPPGLLPPPPPGAPL
	ŀ					LPFPPVSMMPNGPMPVPQMM
						FGLPSLAPLVPPPTLLVPYPVI
						PLPVPIPIPIPIP\PSATPSPPTGS
		İ	İ			ATGRTSFRTPLATPRRRAASQ.
						DTPCPPGTPSRARPSPRTRPPA
						RARP*AWRPRPPSMAGARWW
						*RGAPA\PPGPPGRGRPARLPR
						AAGPAGRRHRPDRGPPSPAAG
						DPPRAA/PRTSRRSASRAPRSA
			İ			PAAAAGTATAARPPPATQAR.
			1			PRRAPRRPRPATSS*TARAAPI
						SGAKSAEPPPEQPPPPPPPAPP
						KLLSPEEPAVMRARVGQGE
21924	52292	Α	22052	762	2049	RSSFSMGADLLACST/SGLGN
		ļ				PY\DLLRGMEPTLGCSFEDLAI
						A*PGGITOE\KINEMRVAPEOO
						MI\ADIHFMIAAGQDLDWIDA
						GATLLHIAGANGYLRAAELLI
						HGVRVDVKDWDGWEPLHAA
						FWGQMQMAELLVSHGAN\LN
			ŀ			RTSMDEMPIDLCEEEEFKVLL
						ELK\HKHDVIMKSQLRHKSSL
			ĺ			RRTSHRQAS/SVGKVVRRTOP
		1				GTGPNL\YRKEYE/GEEAILWO
						SA\AEDQRTSTYNGDIRET\RT
						OENKDPNPRLEK\PVLLSEFPT
				l		IPRGELDMPVENGLRAPVSAY
				1		YALANGDVWK\VHEVPDYSM
						YGNPGVADATPPWSSYKEQS
						OTLLELKRORAAAKLLSHPFL
						THLGSSMARTGESSSEGKAPL
						GGRTSPYSSNGTSVYYTVTSG
		1	İ			PPLLKFKAPIEEMEEKVHGCC
21925	52293	Α-	22053	2	306	SKEAFAIGLGGGSASGKTTVA
	1	Γ.		1		MIIEALDVPWVVLLSMDSFYR
	ł	ļ	1	I		NAFDFDLIISTLKKLKQGKSVI
	1	1				APIYDFTTLSRKMDWKTLYGA
		1				NVIIFEGIMAFADKTL
21926	52294	A	22054	2	379	
	1-3-7	<u>, </u>		I		

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21927	52295	A	22055	352	1227	GDVSPQLSPRQKSGPLCTSCCA
						WPVCSPHSDAQPSCGEGGLWE
		1				KTPFLSVPHRFVPRAALASAHO
		l	Ì			CHPLPRTLSVLKSTPOVRGMHT
						IIRDKETSRDEFIFYSKRLMRLL
		1				EHALSFLPFQDCVVQTPQGQD\
						*RITGVSILRAGETMEPALRAVO
						KDVRIGTILIQTNQLTGEPELHY
		1				LRLPKDISDDHVILMDCTVSTG
						AAAMMAVRVLLDHDVPEDKIF
		ŀ				LL\SLLMAEMGVH\SVAYAFPR
		İ				VKKSSPRRWDKAGQLTFFRIIP
		į.				GIGNFGDRYFGTDAVPDGSDEE
			l			EVAYTG
21928	52296	Α	22056	72	2746	FLVETGLHHVGQAGLKLLTSN
						DPPASASQSTGITGMSHHAQPR
		ľ				QSSYVPPMSLPASAMPVEGVG
						GDALWAGHASGYLGGGOLWA
						TSEYIPLCSSNAESLDRLLPPVG
i						TGRSPRKRTTRQCKSEPPLLRTS
	l	l				KRTIYTAGRPPWYNEHGTQSKI
	Ì					AFAIVPVEQEKEAGQPQAAAA
l	1	ŀ				GWAPLGPICAGSLPVPHRGSPL
		1				PPAGLGGGSASGKTTVARMIIE
				ĺ		ALDVPWGGSSLSMDSFYKVLT
l						EQQQEQAAHNNCNFDHPDAFD
						FDLIISTLKKLKHGKSVKVPIYD
						FTTHSRKKDWKTLYGANVIIFE
						GIMAFADKTLLEGPGPAPCTGP
		1				EDSIIRHIAAPSLHKASSRANYC
		l	l			VLSMGTDPKLATSPIPOCVGAS
						YPKORCPGEVALARPTDOAAS
		1				VQGLVPVSPCPLLASGHSLARC
			İ		1	LHOLLDMKIFVDTDSDIRLVRR
						LRRDISERGRDIEGVIKQYNKFV
		l				KPSFDQYIQPTMRLADIVVPRG
		1				SGNTVGHRP*LLOHVHSOLEEV
						SGVPRPHGLAPSPSTHFSAATTE
		1				SWPVHLIPLRSLSPASS\SSOPAC
ľ						LCQQELGNCSKKLGLGRGRIPL
		ł				SLPKGRHAGGAALASAHQCHP
						LPRTLSVLKST\PQVRGMHTIIR
			ŀ			DKETSRDEFIFYSKRLMRLLIEH
		l				ALSFLPFQDCVVQTPQGQDYA
		1		1		GKCYAGKQITGVSILRAGETME
						PALRAVCKDVRIGTILIQTNQLT
21929	52297	A	22057	61	341	
21930	52298	Α	22058	1	1086	
21931	52299	Α	22059	1	498	
21932	52300	Α	22060	104	3395	
21933	52301	Α	22061	1	1653	

SEO ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				scquence		
21934	52302	Ā	22062	1	1717	
21935	52303	A	22063	172	13375	EDLFKCIMAHAASOLKKNRDL
21/33	32303	ľ	22003	1''2	15575	EINAEEEPEKKRKHRKRSRDRK
1		l				KKSDANASYLRAARAGHLEKA
1			l			LDYIKNGVDINICNONGLNALH
			ŀ	ŀ		LASKEGHVEVVSELLQREANV
		1				DAATKKGNTALHIASLAGQAE
		1				VVKVLVTNGANVNAQSQNGFT
1						PLYMAAOENHLEVVKFLLDNG
ı		l		l		ASOSLATEDGFTPLAVALOOGH
1		1		l		DOVVSLLLENDTKGKVRLPAL
1				l		
ŀ		l		l		HIAARKDDTKAAALLLQNDNN
- 100 1		ļ			200	ADVESKSGFTPLHIAAHYGNI
21936	52304	A	22064	1	360 2577	
21937	52305	A	22065	1		
21938	52306	В	22066	323	3154	
21939	52307	Α	22067	93	791	IRMSYAEKPDEITKDEWMEKL
1						NNLHVQRADMNRLIMNYLVTE
1		l				GFKEAAEKFRMESGIEPSVDLE
		1			ŀ	TLDERIKIREIDT*KGQIQEAIRL
		l				*FNSLHPEL/LWDTNRYLYFHF
		1				ASNSILIPS*FRQ\GKTRGRRWG
	i	l	l			VAQTQLAD\RGEESRECLTEME
	l	ĺ				RTLALLAFDSPEESPFGDLLHT
						MQRQKVWSEVNQAVLDYENR
	İ					ESTPKL\AKLLKLLLWAQNELD
		L				QKKVKYPKMTDLSKGGD
21940	52308	Α	22068	171	426	
21941	52309	Α	22069	5	655	LLQPSVQLPNGTANTSLSFAAV
						SAVSPPPCRTSTATTLPPPMPSF
1						FCVFPSPSMSPSPSEFLSCIASVS
1						RVHSLSSSSSGSSSTASSLNFSAI
1						MGSSSATASWVLSTTSTPPCPS
		1				ALPSSPAQES*SLAASSSAWLSE
						HPSFSSFTLSWASGAFCDEVSLS
		1				FSGSKLSFLSSKGDCGIPFI*TSA
						CPSSLDFSPCFSSSTSKSSSYSEI
		1	l			GKSPQPI

SEQ ID		Met	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X-Unknown, *=Stop codon, /=possible nucleotide
NO:	of peptide sequence	noa	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21942	52310	Α	22070	87	1180	NQASLEVLPSPVDSPDSIPSLPP/ SDNPNRNFSAAPVSCLPSPLPVA
l						ASWPSDPSSCSCPSPEVFRSRAP
						GVPTPASVVYVTVISSAAGTSSP
1						LRANSSTTTFPAAAVFPTNSSC
l						WCQVSSSATSEKEEALFAIVFSS
						AVSEPANTSLSSAAVSAVSPPPC
						RTSTATTLPPPMPSFFCVFPSPS
						MSPSPSEFLSCIASVSRVHSLSSS
	ľ					SSGSSSTASSLNFSAIMGSSSAT
						ASWVLSTTSTPPCPSALPSSPAO
						ES*SLAASSSAWPVAGISPSGAC
						TFPAGSASGAAKAPSPSWRCPS
		ľ				FRALFSLLDSSSLSLCSVSLASL
						SLSPPPSFLRAVMFPATFKSRRC
						FKYGNHCLFSDLLWLEVCAL
21943	52311	A	22071	841	1052	IGEVLLDNVLKSVF*LGSVLPIT
						SLKLLP/SGSKRDPAKMSHSHP
		1			1	AG*VTTGAFPRCGRWRLQPAA
						PPPPS
21944	52312	Α	22072	817	866	HANPVSPPTSWPKE*SWP*CSG
		1				**RTFQSVSFAGGRRRGSWLKP
						PPATAWKSSSCHLPSGVRMAHF
		l	1			CRIPFGPRGRDFADFGTTIKQDF
i		l				RLLGQTSVDRLLQLSQGQAVK
	1					GNQLLPVSLVKRKTTLAPNTQT
						ASPRALADSLMQLARQVSRLES GO
21945	52313	A	22073	2	3238	TTFPAAAVFPTNSSCWCOVSSS
21943	32313	^	22073	ľ	3230	ATSEKEEALFAIVFSSAVSEPAN
1	İ					TSLSSAAVSAVSPPPCRTSTATT
	1					LPPPMPSFFCVFPSPSMSPSPSEF
	1					LSCIASVSRVHSLSSSSSGSSSTA
1	1					SSLNFSAIMGSSSATASWVLSTT
		1				STPPCPSALPSSPAQES*SLAASS
İ						SAWPVAGISPSGACTFPAGSAS
1						GAAKAPSP\$WRCPSFRALFSLL
		1				DSSSLSLCSVSLASLSLSPPPSFL
		1				RAVMFSSDFSNPGVASNMGTT
						VSF
21946	52314	С	22074	88	4374	WEST TO VOLVE DE STATE OF THE S
21947	52315	Α	22075	73	616	VKESLTQKSKRFQNKLINRRG
		1		1	1	WERPRSGASWTQRRALGGRGC VDRWLGII/WPHRIPGARLRGFP
1					1	ALNPTPQTAPSPSPDPRPRPP/CP
		1			1	SDPGPRTEAQCSL\PPPP\PSPRH
		1				LDPGPEPPTPLTQQGQKGTRAP
		1				RPRCPRSQK*ATSGPAVRRREA
					1	RGSASLRTRAPSRDSTHAGIQG
						NVALQP
		L.—		L	I	1

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	eodon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21948	52316	Α	22076	1	421	GTVVMGLPKYFISLSTKFPLSFS
21940	32310	n	22070	ľ	[""	LSADELHIASLFLWPRSKPFGVI
i .						QRSCASQLNALPEVLKNPGDPD
						KMLRFAESPRNLPAGVL/GDIW
	l	l				TDGKGSVSLQEQVYHFPLIRPT
1		i		i		PGS*ARASFLSCG*EEFRRHGLT
						FSSTRE
21949	52317	В	22077	211	2041	· · · · · · · · · · · · · · · · · · ·
21950	52318	Α	22078	200	1525	IMASAHLATSAFCTRSSSSGRSG
						SCKRGSWDSEQALIRAWTTLPS
1		[SLHAADGSFAGGRGIPSKRWS
		1				WRAGQQSSLCLIMQTSSNCGNS
						FQLVSEGASWRGLPHCSCAELQ
1						DSLNFSYHPSGLSLSVRPSSPGN
1						SPKEQPFSQVLRPEPPDPEKLPV
						PPAPPSKRHCRSLSVPVDLSRW
	ļ.					QPVWRPAPSKLWTPIKHRGSGG
						GGGPQVPHQSPPKRVSSLRFLT
		1				SSQCLFSMCPSSQTLQPSFLQPG
						PGP\SSSRPCAASPQSGSWESDA
						ESLSPCPPQRRFSLSPSLGPQAS
						RFLPSARSSPASSPELPLAT*G/L
						SATFPESRSQPCDLDARKTGVK
		l		1		RRHEEDPRRLRPSLDFDKMNQ
		l				KPYSGGLCLQETAREGSSISPP
						WFMACSPPPLSASCSPTGGSSQ
1	(VLSESEKEEEGAVRWGRQALS
						KRTLCQRDFGDLDLNLIEEN
21951	52319	Α	22079	1	117	
21952	52320	Α	22080	3	632	
21953	52321	Α	22081	1	448	MQSETETVHLFIPALSVGAIIGK
	İ	l				QGQHIKQLSRFAGASIKIAPAEA
1		l		1		PDAKVRMVIITGPPEAQFKAQG
		l			1	RIYGKIKEENFVSPKEEVKLEA
1		l	1	1	1	HIRVPSFAAGRVIGKGGKTAST
1		l	1			*QNLCSGMKGKALKAKPSTYN
	1					CVGCGWSPRPRCSLN

SEQ ID NO:	SEQ ID NO: of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21954	52322	A	22082	233	2096	RREDEONKTSWIHPSPWVVLIF RFSLIFFRGVPLTMNKLYIGNL SENAAPSIDLIKSIFKDAKIPVSG PFLVKTGYAFVDCPDESWALIK AIKAFSGKLEHGKPIEVEHSVP KRORIRKLQIRNIPPHLQWEVL DSLLVYGYGVYESCEQVHTDSET AVVNVTYSSKDQARQALDKAL LDKLNGFQLENFTLKVAYIPDE MATQQNPFLQPRGRRGLGGRG SISRQGSPGSVSKQKPCDSPLRL LVPTQFVGAIIGKEGATIRNITK QTQSKIDVHKKENAGAAEKSIT JESTPEGTSAACKSILEIMHKEA QUIKFTEEIPLKLIAHNNFVGRLI GKEGRNLKKIEQDTDTKITISPL QUIKFTEEIPLKLIAHNNFVGRLI GKEGRNLKKIEQDTDTKITISPL QUIKFTEEIPSAMTPPSFOSEQSE SETVHLFIPALSVRALISKQGQH IKQLSRFAGASSKIAPVEAPDAK KRWMMIAGSPEARFKAGGRIY GKIKEENFVSPKEEVKLEHIR GKIKEENFVSPKEEVKLEHIR GKIKEENFVSPKEEVKLEHIR VRSFAAGRVIVFRGGKTVINELQ NLSSAEVVPRDQTPDENDQV VVKITGHFYACQVAQRKIFEEL TOVGOHOOGALOSGPPOSSR

SEO ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nuelcotide
	sequence		09/540,217	eodon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
21955	52323	Α	22083	8	1675	TASCRLAMALHVPKAPGFAQM
						LKEGAKHFSGLEEAVYRNIQAC
					1	KELAQTTRTAYGPKGMNKMVI
	1					NHLEKLFVTNDAATILRELEVQ
						HPAAKMIVMASHMQEQEVGD
						GTNFVLVFAGALLELAEELLRI
						GLSVSEVIEGYEIACRKAHEILP
	l					NLVCCSAKNLRDIDEVSSLLRT
						SIMSKQYGNEVFLAKLIAQAC\
						YLFFLIPAISMLITSEFVK/FLGS
		1				GISSSSVLHGMVFKKETEGDVT
						SVKDAKIAVYSCPFDGMITETK
İ						GTVLIKTAEELMNFSKGEENLM
						DAQVKAIADTGANVVVTGGKV
						ADMALHYANKYNIMLVRLNSK
						WDLRRLCKTVGATALPRLTPP
						VLEEMGHCDSVYLSEVGDTQV
			ĺ			VVFKHEKEDGAISTIVLRGSTD
			1			NLMDDIERAVDDGVNTFKVLT
1						RDKRLVPGGGATEIELAKQITS
			l			YGETCPGLEQYAIKKFAEAFEA
						IPRALAENSGVKANEVISKLYA
	Í					VHQEGNKNVGLDIEAEVPAVK
	ĺ				1	DMLEAGILDTYLGKYWAIKLA
			Ì	i	1	TNAAVTVLRVDQIIMAKPAGGP
						KPPSGK*NWLNFYCR
21956	52324	Α	22084	175	278	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21957	52325	Α	22085	208	1976	GPSPLKEHFSGLEETAYM\NILT
						CMELAQTTRTAYGPNGMNKM
						VINHLEKLLVTNDAATILRELE
		l	l		i	VQHPAAKMIVMASHMQEQEV\
		ŀ				GDGTNF\VLVFAGSFPVNYLKN
			ł		1	L\WRIGLSVSEVIEGYEIACRKA
		ŀ				HEILPNLVC\CSAKNLRDIDEVS
					ĺ	SLLRTS\IMSKQYGLMKVF\LAK
						PYLLQACRIYFSWIPAISMLITSE
		l			ĺ	FVK/FLGSGIRSS*VLAGMVFK
					i	KET\EGDVTSVKDAK\IAVYS\C
		1				PFDGHG*QET*GEQCLDKRTAL
	ŀ			İ	l	KDLDGIFR*GEQENLHG/AATSP
		1				KAIADTGANVVVTGGKVADM
	i	l				ALHYANKYNIMLVRLNSKM/W
	ŀ	1				DLRRLCKT\VGATALPR\LTPSL
		l				SLKEMGHCD\SVYLSEVGDTQ\
		i				VVVFKHEKEDGAISTIVL\RGST
		1				DN/L*LDDIER\AVDDG\VNTFQ
						GSLQRDKRLVP\GGGSNRNLN*
						PKQITS\YGET\CPGLEQYA/ILK
					İ	KFA*GHLKAIPPAHWQKYSGE
		l				NSGVKAQWK*SPKLYAVHQEG
		l				NKNV\GLDIEAEVP/SLVKDML
		l			•	EAGIPRYLTLGKYWAIPLATNA
					i	AVTVLRVDQHTLV*FFFMQIIM
	i					AKPAG/GGPKPPKWGRKTWGD
						DQNGLKIGFNFYCR
21958	52326	Α	22086	1	346	PF*CTPSCLWAF*TARSAQTWS
						APSTRS WCSEWPKSLPSPTWLR
						*FRKVSSYSWSQSWSSPTASTD
						SSRPPHSLGASCHPGHQRSLMP
						PSR*RATSTAYASLSSSHRPNTQ
						PSPS
21959	52327	Α	22087	3	484	RSSQPRVGLCGKSLQP/SSEGSC
						HLQGWCSFPRSVSSIMAEA*RT
						*PRVSSSMHGRGPTQNREATKK
				İ		ASSSETSV/QVTAQEAQAEPGA
1						AGVAG/SPASAPQDSAGACWPA
1	1	1		l		QVL*GGSDRSCRGSHPGNTEPP
						SVLRTGTAPGA*G/SSPFSPTTQS
						GSCCFPFSS
21960	52328	В	22088	181	359	

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nuclcotide location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \-possible nucleotide insertion)
21961	52329	Α	22089	290	2014	STCTPLPTASSQSRPHPPPTPTPP
						APWRSSNGQVPTLT*PGSTLPP
		1				G/SGKSPP/VASGGPARAKPHRA
		1				VHVSPAPSGGSPGDVAELPAAV
		İ		ŀ		AVRA*QAGSGQRLPAPVCVGE
		l				MGTLCPGEPADVHQVVCGLSE
	ļ.	l			1	PRAPHRPGQPQARAHGVPSGQS
						LCPAQPG*DDS/AKVSSYSWSQ
		l			l	SWSSPTASTDSSRPPHSLGASCH
						PGHQRSLMPPSR*RATSTAWRA
						RTASTPRCLGPRPAPWSCASLSS
						SHRPNTQPSPSPTSVRRARLATP
						SSHGWALAPWTPMAPTQPTTW
						TRWGKTVSGRQMNTWRRPWS
						TCARYSGSAKRSSGSSHSPWAP
		l				PRMRMEKSNSPTASWVRTDSS
						LRPWGGTRSSMGCEGLKLSTR
						GTRSCSPSGAMRSPAWSAHSLG
						CRLPSTTDLQDRWRLCVPGMT
				ļ		SSAASVATTSQNLGWPAGTC*A LWGGGRWPATPAA\QAQPALP
		l				GQLPTRWSRCCWPSSWPLCSAS
]		GPSHARCCSPWAMSSTPLP*HC
					1	*PSGGSCTSPEGVSCLQSRLEGF
		l				ATQPHPWAERTWEAPPGGNTV
						ILGLLERGSCSRRGIWRKRNQE
						RKAGRAPAKE
21962	52330	Α	22090	363	487	
21963	52331	Α	22091	234	282	
21964	52332	A	22092	742	990	
21965	52333	A	22093	1	206	LFPAFSDSRECKLMKKLLEA\H
		1				EEQNVD\SYTEA\VKEYDSIFPG
		1				WTQWLTTL\LLRIKKTIQG\DEK
21966	52334	A	22094	128	1062	DLR KPFEAAMYNSGKEAEAMALLA
21900	32334	^	22094	128	1002	
		l				EAER\KV\KNSQSFFSGLFGGSS KIEEACE\IYARAANMFKMAKN
		l	1			WSAAGNAFCQAAQLHLQLQSK
		l				HDAATCFV\DAGNAF\KKADPO
1		l				GGPFNCFDCVAFEIY\TDMGR\F
		l				TIAAKHHISIC*DLLRQKLVDIE
		l				KAIA\HYRALADYYQGEESNS\S
1		l			i	ANKCLLKVAGYA\ALLEQYQK\
1		I	l			AIDIYEQGWGPNAHWTSPPPSK
1		l				Y\SAKDYFF\KAALCHFCMDML
		1				QA\KLAF\QKYEELFPVFSDSRE CKLMKK\LLKA\HGEQNVD\SY
		1	l	1		
		1				TE\SVKGITDSISRL\DQWL\TTH CLRI\KKTIRGD
21967	52335	A	22095	1	1038	ODIG III THOO
2170/	22233	<u>'^</u>	EEU73	l'	11030	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
21968	52336	A	22096	20	2482	RVCSSSASTASOAVMADAWEE
2.,,00	122330	ľ.,	22070	[**		IRRLAADFORAOFAEATORLSE
1		1		ĺ		RNCIEIVNRLIAOKOLEVVHTL
		1		1		DGKEYITPAQISKEMRDELHVR
		l				GGRVNIVDLOQVINVDLIHIEN
		l				RIGDIIKSEKHVQLVLGQL\IDEN
		l		ļ		YL\DRVGQEVNDKLQESGOVTI
	l	1		ŀ		SELCKTYDLPGNFLTQALTQRL
		1				AIIIIGHIDLDNRGVIFTEAFVAR
		1				HRARIRGLESAITRPTAVNSLIS
		1				
		1				KYGFQEQLLYSVLDELVNSGRL
		1			1	RGTVVGGRQDKAVFVPDIYSR
		1				TQSTWVDSFFRQNGYLEFDALS
		1			i	RLGIPDAVSYIKKRYKTTQLLFL
		[KAACVGQGLVDQVEASVEEAI
1		1				SSGTWVDIAPLLPTSLSVEDAAI
		1				LLQQVMRAFSKQASTVVFSDT
1		ł				VVVSEKF\INDCTELFRELMHQ
		1				KAEKEMKNNPVHLITEEDLKQI
		1				STLESVSTSKKDKKDERRRKAT
		1			1	EGSGSMRGGGGGNAREYKIKK
					ľ	VKKKGRKDDDSDDESQSSHTG
		1				KKKPEISFMFQDEIEDFLRKHIQ
					ľ	DAPEEFISELAEYLIKPLNKTYL
					ľ	EVVRSVFMSSTTSASGTGRKRT
		1			1	IKDLQE\EVSNLYNNIRLFGKRG
						WKFF\ADDTQGWLLPKHLLEV
		1				QCVLISLNLI\FNF\LGFGILMDG
						Q*DDPGRPFTS\ELRKEI\LSKLS
		l				EETKVALTKLHNSLNEKSIEDFI
	l	l				FC\LDSAAEACDIMVKRGDKKR
				1		ERQILFQHRQALAEQLKVTEDP
21969	52337	A	22097	1	711	VLSLFDGIATGYLVLKELGIKV
		1				GKYVASEVCEESIAVGTVKHEG
		l				NIKYVNDVRNITKKNIEEWGPF
		1				DLVIGGSPCNDLSNVNPARKGL
		1				YEGTGRLFFEFYHLLNYSRPKE
		1	l	l		GDDRPFFWMFENVVAMKVGD
				1	l	KRDISRFLECNPVMIDAIKVSAA
		l	ĺ	1	l	HRARYFWGNLPGMNRPVIASK
	1	l		1		NDKLELODCLEYNRIA\RIFGFP
		1		l		VHYTDVSNMGRGAROKLLGRS
			l	1		WSVPVIRHLFAPLKDYFAC
		L		L	l	WSVI VIKILLI AFLKDITAC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	coden for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		l		sequence		
21970	52338	Α	22098	406	2885	GRGILVNGACSDQSSDSPPILEA
						IRTPEIRGGWAVGTGVVSGADI
		l		l		VSGHCRQWSLLSLFQGSGTLSL
				l		AFITWAFRETRTRSESPAVSSHT
						SSQSTCGQHST*HT*HARGRQG
						RNHVDESPVEFPATRVGSPDAQ
			l	l		TPARSL*LGDMPHSLTIDLTDDT
			i			EDTHGTPQSSSTPYARLAQDSQ
			İ	l		QGGMESPQVEADSGDGDSSEY
				l		Q\DGREGLLPRV/I*GP*TRGKTS
						YAMVVSWKATSKRQAMSGMR
			ĺ	l		WVQWFGDGKFSEVSPGKGKGF
		l				CRPEAVPACLPLNKLVSYRKA
						MYHALEVTWDEGMRAKP*EQ
						G*SKRPTELEDQLKPMLEWAH
						GGFKPTGIEGLKPNNTQPGGNE
						SPWQHPLPLLVGPLLGRVRLKT
		1				NCYNNGKDRGDEDQSRGDCW
		1				VPGIMGQMGGGRCRSGIDVIPL
		l				TPWLGL*ACLVCQTCRVSPPTT
	1					ALDLPPLASSNSPLLPCEGRELL
	i					LCSNTSCCR*ALGPVGWMWVG
	1	ŀ				PQGSYVPAV*EPWSCYMCLPQ
						RCHGVLRRRKDWNVRLQAFFT
1	1					SDTGLEYVSHRLPPLPPQIPGAQ
	1					GVGKLWFDGIATGEFGEHLETL
1	1					LSCHNRVARESEESIAVGTVKH
			1			EGNIKYVNDVRNITKKNVRAV
		ĺ				CTLRASSLPVFFPSPPHPNPARK
		1			1	GLYGEHPSLWQSLESLCHLTTG
		l	1		1	PGAGNK/RGLGWRFQMNPALS
		1				SFLST\GVQRDISRFLEVRESGD
21971	52339	Α	22099	729	896	

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	sequence	nou	09/540.217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	Sequence		03/240(21)	sequence	or peptide sequence	
21972	52340	IA	22100	li i	2672	I MAYKOGFLKVMVHFRKAEVT
		1				GKNHTSVHEGYTLISPAAOVLT
	1	1			\	ISCKKSILLOEQVSFKDVCVDFT
		1				QEEWYLLDPAQKILYRDVILEN
		1				YSNLVSVGYCITKP\EVIFKI/ES
ļ		1				KGEEPW\VLEKGIPOAOCPPOK
		1				RKWKV/EMNVVREQPRKMKM
		1				TIFGEPPIPTTNKTVSVENGDRG
1						SKTFNLGTDPVSLIKLSL*NM*L
		1				SEMNLKNISGLIITKKNCSRKK
				i		PDEFNVCEKLLLDIRHEKIPIGE
Į.						KSYKYDQKRNAINYHQDLSQP
		l				SFGQSFEYSKNGQGFHDEAAFF
						TNKRSQIGETVCKYNECGRTFI
						ESLKLNISQRPHLEMEPYGCSIC
						GKSFCMNLRFGHQRALTKDNP
]		l				YEYNEYGEIFCDNSAFIIHQGAY
		l				TRKILREYKVSDKTWEKSALLK
1		l				HQIVHMGGKSYDYNENGSNFS
l		l				KKSHLTQLRRAHTGEKTFECGE
		1				CGKTFWEKSNLTQHQRTHTGE
	1	l				KP\L*CTEC\GKAFCQKPHLTNH
						QRTHTGEKPYECKQCGKTFCV
						KSNLTEHQRTHTGEKPYECNA
						CGKSFCHRSALTVHQRTHTGE
		1				KPFICNECGKSFCVKSNLIVHQR
	1					THTGEKPYKCNECGKTFCEKSA
	1					LTKHQRTHTGEKPYECNACGK
					1	TFSQRSVLTKYQF\IHTRVKAL*
		1			1	TS*MLEAFIHL/SKLVIQFQKRRS
					1	EKAKECQKFVE\KTSCLNM*KL
		<u> </u>				SRKIKTFH*KICKLTVKNKGNK
21973	52341	Α	22101	1	2817	1

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
21974	52342	Α	22102	474	2766	NQTALYNQAQLLHGSRFANMS
						AEGYQYRALYDYKKEREEDID
	1		İ			LHLGDILTVNKGSLVALGFSDG
				ł		QEARPEEIGWLNGYNETTGERG
						DFPGTYVEYIGRKKISPPTPKPR
				1		PPRPLPVAPGSSKTEADVEQQA
	ľ					LTLPDLAEQFAPPDIAPPLLIKL
						VEAIEKKGLECSTLYRTQSSSNL
						AELRQLLDCDTPSVDLEMIDVH
			1			VLADAFKRYLLDLPNPVIPAAV
		İ				YSEMISLAPEVQSSEEYIQLLKK
						LIRSPSIPHQYWLTLQYLLKHFF
						KLSQTSSKNLLNARVLSEIFSPM
						LFRFSAASSDNTENLIKVIEILIS
	l					TEWNERQPAPALPPKPPKPTTV
						ANNGMNNNMSLQNAEWYWG
						DISREEVNEKLRDS\ADGT\FLV
		l				RDASTKMHGDYTLTLRKGGNN
					1	KLINKIFHRDGKYGFSDPLTFSSV
		1				VELINHYRNESLAQYNPKLDVK
l						LLYPVSKYQQDQVVKEDNIEA
l				1		VGKKLHEYNTQFQEKS\REYDR
		l			l	LY\EEYTRTSQEIQMKRPAIEAF
				l		NETIKIFE\EQCQTQERYSKGNY
1		1				HLEKFKP*RAMEKEIQRIMHNY
		ĺ				DKLKSRISEIIDSRRRLE\EDLKK
				1		Q\AAEYREIDKRMNSIKPDLIQL
				l		RKTRDQYLMWLTQKGVRQKK
	l		l	I		LNEWL\GNENTEDHYSLV\EDD
					1	EDLPHHDE\KTWECLGSS/RTRN
						QSLKNLFARGSRDGTFLVRESS
	l	1		L		KQGCYACSVVVDGEVKHCVI\
21975	52343	Α	22103	3	392	
21976	52344	A	22104	168	404	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21977	52345	A	22105	49	1760	YQSPAWQSHWHQKLQVRKEN MAKRVAIVGAGVSGLASIKCCI EEGLEPTCFERSDDLGGLWRFT EHVEEGRASLYKSVVSNSCKE LEVLKMYANHFDLLKHIQFKT KVCSVTKCSDSAVSGQWEVVT MHEEKQESAIFDAVMVCTGFL TNPYLPLDSFPGINAFKQQYFH MEEKQESAIFDAVMVCTGFL TNPYLPLDSFPGINAFKQQYFH GYKHPDIFCDKRVLVIGMEN GTDIAVEASHLAEKVFLSTTGG GWISRIFDSGYPWDMYFMTR FQMILRNSLPPITYTWLMERKI NNWLNHANYGLIPEORTQLEE FVLNDELPGRITGKVFIRPSIKE VKENSVIFNNTSKEEPIDIIVFAI GYTFAFPFLDESVVKVEDQQAI SLYKIPFAHQFFTLAIGH IKPLGLP*LPTGETQVRWAVSG S*KVLTRLPPFSVMIEEINARKE MFSWFGLCYCKALQSPVITYI DELLTYINTKPNLFSLWILTD FIGGLWYFGPCSPYPGFLDFT FGGKMGKGAQEIALHGPQWDP NISRVHSKVRSCYKESIFFFCK
21978	52346	A	22106	262	969	EWSSVRRSLVEKRALRRPHPQG LEFRMKTILISNQTVDIPENVOI TLKGRTGIREGPPEGTLERIDFN HINIVELSLLWEEKKRGSRVDK WWGNOKRNWPTRSD*FVSHV QNMIKGWLHWGFRLQG*GPV YAHFPHQPLLSRKNGVSWLKIF KFLWGEKYNRVSG*RFGCLL VSVSQCPRKDENPLKGNDIEL VSK/SSAGF*FQQATTS*KTRNI RKFLNDGICLLEKGLFRQA
21979	52347	В	22107	211	457	
21980	52348 52349	A C	22108	167	367	KEAENQCLDISSPGMSGHRQGF DSLEHDELREIFNDLSSSSEDED ETQHQDEK*LDISSPGMSGHRQ GHDSLEHDELREIFNDLSSSSED EDETQHQDEK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
21982	52350	A	22110	754	1950	VKGKDDAPHELGE/SOFILRLPP
		l				VEYASTVRRAVQSGHVNLKDRL
		l				TVELHPDGRHGIVRVDRVPLAS
		l				KLVD\LP\CVMEEPENPLLKKPF
						SKTAHGOMLVSTVDGDLFPPV
		l				EEPSCLALDPKAS\KKKDKDKE
						KK\FIW\NHGITLPLKNVRKRRF
		1				RKTAKKEIYLNLPDV*KKELKR
		l		l		\LLEVPDA\EAVSTRWGIICPKV
		l				ETKG\AENQG\LDISSPGMSG\H
						R\QGH\DSLEHDELREIFNDLSSS
		l				SEDEDETOHODEEDINIIDTEED
						LERQLQDKLNESDEQHQENEG
	ł			l		TNOLVMGIOKOIDNMKGKL\OE
	1	l				TODRAK\ROEDLIM\KVGNLGS
	1			ľ		PRTDFQAVLGMSFNQREDREK
	1	l		l		EQL\SSLQEELESLLREVKKNLM
	1	l				FNFQSFRLVQHLGKFFGLLLLL
						GYLRPCSS
21983	52351	В	22111	227	355	
21984	52352	Α	22112	121	486	WGPALCIGGYLASFLASVHLMP
		l				LR*GEMLSP*GLSFSFRFPGFKE
				i	1	VRLVPGRHDIAFVEFDNEVQAG
						AARDALQGFKITLSPLKVSPPL
						GAFLEPCVSEWSPHSIVPRVCPQ
						TLHLALLGRN
21985	52353	В	22113	8	165	
21986	52354	Α	22114	1	218	
21987	52355	A	22115	566	1412	TLHSMAVPETRPNHTIYIN\NLN
						EKIKKDELKKSLLRPSFF/SSLG
		l				QILGYSWGIHGRP*RMEGARPF
		l		l		SIFKEVSSGTNALR\SMQGFPFY\
						DKPMRIPVLPRTRLQDIHLPRIE
						RAPFIGSRDPKAGERGRPKSQG
						DPGPPKEGLCKGGGSHPRGGG
		l		i		LSRGPGPGHCPPMT\KAPRI\MH
			1	l	1	HMPG\QPPLTAAPLDMIPPPG\L
		ı	l	[1	APGQIPPG\AMPP\QQLMPGQMP
				l		P\AQ\PLWRNPP\NHILF\LTNLP\
			1	l	1	EET\NELMLSHAFSNQVPPGFKE
l	1	ı	1	l	1	GPVWVPRAGNEHRPSVEV

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21988	52356	A	22116	2	1176	FGELSTFDAVRGPPRRCPPDAD PRAPGLALPGRLLVSNSRRTHS RESRAPPAPHLLGARAGMAPPO
						VLA\FGLLLAAATATFAAAQEE CVCENYK\LAVNCFCE*INRQC
						Q\CTSVGAQNTVICSKLGCQMF W*MKAEMNGSKLGRRAKPEG GLO\NNDGLYDPDCDESGLFKA
						KQCNG\TSTC\WCVNT\AGVRRT HKDTEVTCSERVRTYWIIIELKH
						KAREKPY\DSKS\LRT*LQKEITT RYTLGSKFLSRVFWYENNVITI\ DLFONFFSKKLRNDVDIADVAY
						YF\EKDVK\GESLFH\SKKMDLT VNGEQLDLDLLIPGQTL\IYNVD
						EKAP*/DFSMQGLK\AGVIAVIV VVVITSCCWKLLVLVISQKGRE WOKYEKAEIKGDGCRCHRELO
21989	52357	A	22117	3	257	AKALQWPLGRGGGGTPRGSRP PGPH*SERGAVQPRSLPRQAA* RVPSGPADGQ*DGHGAADPGS
21990	52358	A	22118	1467	1748	RQRCKPWSEAADLHSECYSS SEYNCCEIGCPQPSWGPWGRG\ TDVRQGQVGEEEVHG/ECGGG
						SQG*RPG**AGSPAP*PGTWTGT AQQGPAAVLDPLRV*GGGIERH PLHSVGL
21991	52359	В	22119	1	1461	
21992	52360	Α	22120	29	417	

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of tast	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	coden for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	detetion, \=possible nucleotide insertion)
				sequence		
21993	52361	Α	22121	616	1671	TATCKPWSMRTTISSSQPQKLT
		1				NSVKLPPSAKCISVSRIGGFLVS
		l				LTSRMKLRTLLICSKLLPSGGV
	l	1				RGLAGSGVKLQTFTVSVTALK
		1				AACLELFTSPGGFMVLLASEVK
	İ	1			l .	LOTFSAGDSDCLGQGRRAQRRI
	ı					GPGVSAVGASGQQRAPESNERI
	l					AEEPTPAPMPSSAPRGPTEPALR
						RPSSADRPLAPGAGRAPGAMAS
1		1				PSGSSKATGKPRGRDGRPRREE
i	l	1				DDVTPEEKRLRLLLEEGSAQPD
	ł					DGEDAPRPGRKETGTQTGGDG
l						RGAARAALLAVGGIRAPLPSLG
						VAPGC\PMSKPPRRHPGATPR/P
	ļ	1				GQGGSDAAHGEEGCPCCSSAV
		1				ATCLGAGLLP\PGRGASSPSSGC
	ł	1			i	ALPSSSNSRSLFSSGVTSSSSLLG
	i	l		İ		RPSLPRGLPVALELPEGLAMAP
						GALPAPGASGLSAELGRLSAGS
	i	1				VGPRGA\G*ASGRAWAPPRAAR
	İ					CSLAPSAG\PRAHRGHAGPWPA
	l					LSSP/GPGPSSRCPLPRRSAASLL
						KPAR/AINPPGEVNNSRHAALR
			l			AVTLTVKVCSFTPEPARPRTPPE
	l					GRSFEHIRRVRSFILEVSETKNP
	Ì					PILDTLIHLALGGSFTLFVNFCG
	l					WDELIVVLIDQGLHVAV
21994	52362	A	22122	892	1931	CTWNENQRMEIVWEVLFLLQA
	ŀ					NFIVCISAQQNSPKIHEGWWAY
	l	l				KEVVQG\NFVPVPSFWGLVNSA
						WNLCSVGKRQSPVNIETSHMIF
						DPFLTPLRINTGGRKVSGTMYN
			Ì			TGRHVSLRLDKEHLVNISGGPM
		1				TYSHRLEEIRLHFGSEDSQGSEH
	l					LLNGQAFSGEVQLIHYNHELYT
						NVTEAAKSPNGLVVVSIFIKVS
1	l	l				DSSNPFLNRML\NRDTF\TRN*H
						FKNDGIFYYRGL*YRGELLFQG
1	Į.					TSSFITYDGVDGLFPTPGY/SRT
	1	1	l			ASWIIMNKPV*YNPGCKMHSLR
	1	1	l			LL\SQEPAISRSFLSMSDNFRPV
1	l	1	1			QPLNNRCIRTNINFSLQGKDCP
		1				NNRAQKLQYRVNEWLLK
21995	52363	Α	22123	17	417	
21996	52364	Α	22124	63	449	CHPALLGPTGHGAWPRHRAPD
l	1	1	1			TLRASAGGRDP*GRRSAPA*ER
1	1	1	1	I		QP*PPVYCAGISQYD*TALSSCL
l		Ì				RRSVRTSTDFRSLLDKSNYSVF
l			1	I		LCFLQLAGRRYFAMWTRPEFFL
		1		I		ARDPRSLSWGLEWRPLSL
21997	52365	В	22125	1	1131	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nuclcotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
21998	52366	A	22126	359	548	
21999	52367	Α	22127	755	1031	TRGVLEHMALGKGTLKDPSGP
						AP*NHPE\VSHAASTQEAAGPP
	l					AGRPGHGSGCLALPASSTSLEP
		i				GLRQDHDLLLQLLWREQEVFH
						DVIQLL
22000	52368	Α	22128	1	277	
22001	52369	Α	22129	1	933	
22002	52370	Α	22130	339	863	PSPVPARPVCPSSAPISMRPPCP
		l				WREAEAADLRPEAPGHRDGED
						VGFLSSGQLSGLCSKRKLQPTR
			İ	l		AALTLTPSAVNKIK\HFLKNNPE
		l				HVGVKVGVRTRGCNGLSYTLE
						YTKTKGDSDEEVIQDGVRVFIE
1			1	l		K\KAQLTLL*TEMDYVEDKLSK
1				ŀ		GVGV\ITPNIKGTCGCGESFNI
22003	5237I	Α	22131	16	419	QIQHRPWP/SRSYTNVLGRKVV
		1				YLPSFFTYAKYIVQVDGKIGLF
1				l		RGLSPLMMSNALYTATRCFMK
		1				KDLPPDEIEQVSNKDDMTTSLN
				ľ		KVANDTSYEMMMHCVSRMLP
	ĺ	l		ł		HPLHVISMRCMDQIVGREAKYS
						GVLSSIG
22004	52372	A	22132	3	441	RSYTKFVMGIAVSMLTYPFLLV
		l				GDLMAVNNCGLQAGLPPYSPV
İ						FKSWIHCWKYLSVQ\GASTGRR
		i	l			RPFLFFATSFS*NGFVDASLHAK
		1				GQLFRGSS\CFSAGCHQDHALP
						WSNLNHLKNTVSTWPPWVRPD
						HLGTPARRLQPNNNQM
22005	52373	Α	22133	1	1232	MAGAGAGAGARGGAAAGVEA
		1				RARDPPPAHRAHPRHPRPAAQP
			1			SARRMDGGSGGLGSGDNAPTT
						EALFVALGAGVTALSHPLLYV
				İ		KLLIQVGHEPMPPTLGTNVLGR
						KVLYLPSFFTYAKYIGQRDG\RE
				1		GLVPRLSPRLMSNALSTVTRGS
						MKKVFPPDEIEQVSNKDDYEDF
				l		P*RKLLKETSYEMMMQCVSRM
						L\AHPLHVISMRCMVQFVGREA
						KYSGVLSSIGKIFKEEGLLGFFV
				1	1	GLIPHLLGDVVFLWGCNLLAHF
				1		INAYLVDDSVSDTPGGIGKRPE
				1		SRFP\FSQALALRSYTKFVMGIA
l	1		1	l		VSMLTYPFLLVGDLMAVNNCG
			1	l		LQAGVPP\YSPVFKSWIHCWKY
				1	1	LSVQGQ\LFRGSS\CFSAGCHQD
1		ĺ		1		HALPW\SNLNHLKNTV\STWPP
		l		1		WV\RPDHLGTTARRFQPNNNQ

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
22006	52374	A	22134	3	1156	PRHPRPAAQPSARRMDGGSGG
						LGSGDNAPTTEALFVALGAGV
ĺ		l				TALSHPLLYVKLLIOVGHEPMP
			l			PTLGTNVLGRKVLYLPSFFTYA
						KYIVQVDGKIGLFRGLSPRLMS
			Ì	1		NALSTVTRGSMKKVFPPDEIEO
						VSNKDDMKTSLKKVVKETSYE
						MMMQCVSRMLAHPLHVISMR
						CMVQFVGREAKYSGVLSSIGKI
						FKE\EGLLGFFVGL\IPHLLG\DV
						VFLWG\C\NLLAHF\INAYLVDD
					ľ	SFS\OAL\AIRSYTKFVMGIAVS
			ł			MLTYPFLLVGDLMAVNNCGAA
						SWGSPPYSPVFKSW\IHCWKYL
		1				SVQGQLFRGSS\CFSAGC\HQDH
		1	ŀ			GFALGVN*NHLKNT\VSTWPPW
						VRPDHLGTPARRLQPNNNQMC
		l				SSPAG\FQFHICHVSV\QMWG
22007	52375	Α	22135	2	425	
22008	52376	Α	22136	2	430	
22009	52377	Α	22137	2	406	
22010	52378	Α	22138	1	297	
22011	52379	Α	22139	1	111	
22012	52380	A	22140	1	398	
22013	52381	Α	22141	1	409	
22014	52382	A	22142	500	1234	GPNCRPPSCSAVEGAWPRPQGA
						GARAAIALSQKLELGGVGLVQ
		1				VQCPSLAPSSTSSILFLCLLPHPP
						PHRCLQENTTCLLGVEGVLLLQ
						SPFPLLKTDHSSLPALAPTQSQA
						ALWGFADFGFMFESSPGGNLG
						WEPDIKLTDEMVMIMGGKMEA TPFKWFMEMCVRGYLAVRPY
						MDVVVSLVTIMLDTGLPCFRG/
	İ					QIKFLKHRFSPNMTEREA*NFIM
						KVIQSCFLSNRSRTYNMIQYYQ NDIPY
22015	52383	A	22143	2	6378	SAGPAGPLQVQKLLCMCPVDF
22013	32363	^	22143	 	0378	HGIFQLDERRRDAVIALGIFLIES
						DLOHKDCVGPYLLRLLKGLPK
						VYWVEESTARKGRGALPVAES
			1			FSFCLVTLLSDVAYRDPSLRDEI
						LEVLLQVLHVLLGMCQALEIQ
						DKEYLCKYAIPCLIGISRAFGRY
		ĺ				SNMEESLLSKLFPKIPPHSLRVL
						EELEGVRRRSFNDFRSILPSNLL
						TVCQEGTLKRKTSSVSSISQVSP
						ERGMPPPSSPGGSAFHYFEASC
						LPDGTALEPE
	L	L		L		PLOGIATELE

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, "=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22016	52384	A	22144	3	398	IHKAASGVPQQPDAGCPGTGEG QGGQRCGQ*AGP/AQAPC*SRA GTR/PTCFFVLTSLRSWYRVDSP NKLMNPLVAGFFG/CHCGSGQC LRKCSSARDRAPDARPGLDLAQ GRCCHCNKACLHASETGVGTT SVL
22017	52385	Α	22145	344	498	
22018	52386	A	22146	344	406	FLADQEDPLDPRAGCR*ETTAP RIPSKGPSSTEPLHASTAVPPCQ GELGCSRASPASEKI/DLAGRLIS PCNSDTVSPGLGGFMTCAQSLI SRRPGGPIGPPGWLOGPHLG
22019	52387	A	22147	3	2058	LSGSGAPPEOGAQAAAAAE EEREVYRVRVRALAVSRPSRR GKRSFAAVDEOTISLDVLQHILI RAFDLSG*VGRCWGLLGRDRL GQEVYLSLLSDWDLSTAFATAS KPYLQLRYDIRFSEDSEYSVPFG AISPKDVIGSDVLLAERRSSLIT AALPFTGSLIT.OYPSGTLGTQQ VLSWSYGEDVKPFKPPLSDAEF HTYLNHEGQLSRPEELR.RIYH GGVEPSLRKVSILSHLLVYPDG LTGRERMDYMKRKSREYEQLK SEWAQRANPEOLEFIRSTVLL VRTDRAHPYYAGPEOPHILR ALHDLLTTYATTHPQVSYGOM MSDLASPILAVMDHEGHAFVC FGGIMKRLAANFHPDGRAMAT KFAHLKLLLELKREF AEDDALRMLETWSSLPPDPPE HEVELVGPPSQVADTGFGDHR GWPYRQRHMLRPAGGGGSTF EDAVPHLATASQGFGGGRLL RQASLDGLQQLRDNMGSRDP LVQLFPALSSKGLSEGNL YPDPLLSSFSHPDSPSSSSPPSTQ EASPTGDMAVGSPLMQEVGSP KOPGKSPPULFRICALLELRERDHIM VPDPLLSSFSHPDSPSSSSPPSTQ EASPTGDMAVGSPLMQEVGSP KOPGKSLDFWFMLFPAGGGGSTL KOPGKSTAFT KARLALLLELRERDHIM VPDPLLSSFSHPDSPSSSSPPSTQ EASPTGDMAVGSPLMQEVGSP KOPGKSLPPYPMGLPPPQEFG RGMPFMLFLCALLLERRDHIM KNOLDYNELAMHEPDRLVRKHH LGRVLRRARALFADVLQSEVW
22020	52388	Ä	22148	130	387	
22021	52389	A	22149	1	279	

SEQ ID NO:	of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide inscriion)
22022	52390	A	22150	105	499	DMSDTSESGAGLTRFQAEASEK DSSSMMQTLLTV/TPEAREAPA TQASSTTQLTDTQVLAAENKSL AADTKKQNADPQAVTMPATET KKVSHVADTKVNTKAQETEAA PSQAPADEPEPSAAAQSQENQ DTR
22023	52391	A	22151	285	984	TSLSCYFSRTKOSPALGLLMVL LSHIFMINGNRSSEAVIWEVLRKL GLRPCYDWALSALAVRVVL.W GERTVLGLHQSGGLVERVGCW TO*ANQGSDLGG*WNOFPELS APSLSVSWASMELPSCAGNLFS ILEMPLFTSGKCHSLE*P1YFFIFF SFSSLRIHHSLFGDVKKLITDEF VKOKYLDYARVPNSNPPEVEFF WGLRSYYETSKMKVLKFGCKV QKKDPKEWAAQVRE
22024	52392	Α	22152	259	578	
22025	52393	A	22153	296	2242	IMMSDTSESGAGLTREQALEAS KOSSSMMGUTLTVTQNVEVEP TPKASKALLEVSEDVKVSKASG VSKATEVSKTPEAREAPATQAS SYLLSLTDITOVLAALENKSLAA *HOGNRMVDPQAC*QLPATEIT KKVSHVADVGRENVKGFREL LAPS\QAPADEPEPSAAAQFSG RERVFGFKVKAKKARKVKHLD GEKDGSSDQSAGSTTGGRRV SKALMASMARRASRGPIAFWA KKALKALWARRAFALPLEI TLKPRRAKGLRRRVAKFQSSQ EPEAPPRGILALLQGRANDLV KYLLAKDOTKIPIKRSDMLKDII KEYTDVYPEIRERAGYSLEKVF DAGILGTTKDSPKLGLLMVLLS GILRPGIHHSSLLGDVKKLITDE FVKQKYLGLCOKVPNSNPPEY EFLLGACALT-*EPAKMKSPQS FACK/VYQKEGSPRNWGSSVPP REGRAGGREDADFEGCKLRITLE EKAKGPRLEAPNOPLGLGSGE MLAGPCNWDEAUIGNPWAK RIPREGEKLKAKAPKRSGSAS TGA\STSTNNSASIASASSTSGGF SAAASLTATLINNSASIASASTSGGF SAAASLTATLINNSASIASASTSGGF SAAASLTATLINNSASIASASTSGGF SAAASLTATLINNSASIASASTSGGF SAAASLTATLINNSASIASASTSGGF SAAASLTATLINNSASIASASTSGGF SAAASLTATLINNSASIASASSTSGGF SAAASLTATLINNSASIASASTSGGF SAAASLTATLINNSASIASASTSGGF SAAASLTATLINNSASIASASTSGGF SAAASLTATLINNSASIASASTSGGF SAAASLTATLINNSASIASASTSGGF SAAASLTATLINNSASIASASTSGGF SAAASLTATLINNSASIASASTSGGF SAAASLTATLINNSASIASASTSGGF SAAASLTATLINNSASIASASTSTNSAS

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
22026	52394	Α	22154	1	487	EFRQPSFRRSIWP/YFLLTQA/VR
						TGNPAKFNQVLDQFGEKFQAD
		İ	Ì			GTYTLIIRLRHNVIKTGVRMISL
						SYSRISLADIAQKLQLDSPEDAE
1		ĺ			i	FIVAKAIRDGVIEASINHEKGYV
			l			QSKEMIDIYSTREPQLAFHQEG
l						KDSSVYSFREWGTRTGMSWLV
						CFLKLRL
22027	52395	A	22155	2	429	
22028	52396	Α	22156	1	328	GTRFQADGTYTLIIRLRHNVIKT
						GV/RSGSEGLLEEQKPGQGKPS
1		l				GYPWASEGSVSTLPTPSLLPLPG
1			ŀ			VRMISLSYSRISLADIAQKLQLD
						SPEDADFIVAKVGLVQEPQ
22029	52397	Α	22157	138	2206	AVTPGAMKQEGSARRRGADK
		l				AKPPPGGGEQEPP\PAPAPQDVE
			ļ			MKEEAATGGGSTGEADG\KTA
		1	l		İ	AAAA*SNFPAKSWDYSHLWKD
						IKEHRETS*RKAVSGKEPRFVLR
						ALRMLPSTSRRLNHYVLYKAV/
			ĺ			QGAFFTSNNAT\RDFLLPFLERA
1					i	HGTQEADLLFRPRTGKAAGT\P
1						LLPEVEAYLQLLVVIFMMNSKR
						YKEAQKISDDLMQKISTQNRRA
		l				L\DLV\AAKCYYYHARVYEFL\D
		l				KL\DVVRISLHG\RLRTATLRHD
l					1	ADGQAT\LLNLLLRNYL\HYKL
						VPTSLRSLVSKSVFP/EEQAQQP
						NEWGQVTSYLHQGRIKAIQ\LE
						YSEARRTMTNALRKAP\OHTAV
			ĺ			GFKQTVHKLLIVVELLLGEIPDR
						LOFROPSLKRSLMPYFLLTOAV
						RTGN\LPRFTQVLDQFGEKFQA
						DGTYTLIIRLRHNVIKTGVRMIS
						LSYSRI\SLADIAQKL\QLDSPE\D
		l				AEFI\VAKAIR\DGVIEASIN\HEK
			ŀ		}	AYVQ\SKEMIDIYST\RDPQLAL
						H\QRISFCLDIHNMVWSRPMRV
			İ			PSPKSYKOGSWSLORNGRERR
						NKQDLE\FPKEMA\EDDDDSFPL
1			l		1	KLGGWGGVRGKWGTROLCPP
			l			WGYPLPQGTCPPFSPHTAHML
1			l			AFRGRGWGCWEPATPDLPPGL
1			l			LPSRVTLLVQPGRRVGRQPPRG
			l			RVLASSVADRRGRISLCTPFRE
1			1			WGTRTWDVYGLLCFLKLRL
22030	52398	Α	22158	86	348	
	102070	٠.,	1-2.00		I -	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nuclcotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \-possible nucleotide insertion)
22031	52399	Α	22159	3	356	LPLSMDVRAPPAVMAAVDQAL
l	1					KEFGRIDILINCSSSSCGLPFCR\
						AAGNFLCPAGALSFNAFKTVM
						DIDTSGTFNVSRVLYEKFFRDH
						GGVIINITATLGNRGQALQVHA
	l					G\SAKAAVD
22032	52400	A	22160	3	255	GPWSCHAGA*GPGATQVSEDQ
						GLREPHGSRDLCSLG*KH*GAPI
	1					CAWPMSWDGPPAAHKVRPSVS
						APPVCINTLCYNLASAFS
22033	52401	Α	22161	3	360	
22034	52402	Α	22162	240	1356	AGNPSGIPGTRVHRGQRGGFAS
	1					TWSGRGAGSTPLQNSPPRQPHR
						EKEFLRQVLEGGSHTGVFRPSA
						ARLVPNPHSLVSPDLL*LVSRA
						T\PHIFLAARKLAGATGRRCLPL
						SMDVRAPPAVMAAVDQALKEF
		1				GRIDILINCESVLSEVGGFSQAP
						AAPAVSHSAGAAGNFLCPAGA
		l				LSFNAFKTVMDIDTSGTFNVSR
						VLYEKFFRDHGGVIVNITATLG
						NRGQALQVHAGSAKAAVDAM
						TRHLAVEWGPQNIRVNSLAPGP
						ISGTEGLRRLDASQGMSWGRG
l	l		1			EGAGSWGSRGLSLLLPS\GGPQ
1					1	ASLSTKVTASPLQRLGNKTEIA
l						IISVLYLASPLASYVTGAVLVA
		_	2014		1.00	DGGAWLTFPNGVKGLPDFASF
22035	52403	В	22163	214	1647	

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SEO ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown.
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
20026		<u> </u>	00164	1.000	lioro	L CALIFORNIA DE LA CALI
22036	52404	Α	22164	1765	1910	LSHPHSRTPPNRTPRRTRIPQRP
İ					\	AVMYSPLCLTQDEFHPFIEALLP
ŀ				i		HVRAFAYTWFNLRARKRKYFK
						KHEKRMSKEEERAVKDELLSE
						KPEVKQKWASRLLAKLRKDIR
l						PEYREDFVLTVTGKKPPCCVLS
						NPDQKGKMRRIDCLRQADKV
					1	WRLDLVMVILFKGIP\LESTDGE
l				ŀ		RLVKSPQCSNPGLCVQPHHIGV
l				ŀ		SVKELDLYL\AYFVHAGRISSQS
1						ESP\SQ\PSDADIKDQPENGTFGA
			1			SRDSFVT\SGVF\SVT*A*LRVSQ
	1			ĺ		TPI\AAG\TGPNFSLSDLESSSYY
	1					SMSPGAMRRSLPSTSSTSSTKRL
						KSVEDEMDSPG\EEPFYTGQAR
1						SPGKWTVSPSGLSVSWSQGMPS
						PTTLKKSEKSGFQQPLPFHRPPS
						LGNGFHTAFTDLFIYQGPQSKF
1						HIATPSIL\HFPRHSPFFQQPGPY
						FSHPAIRYHPQETLKEFVQLVCP
						DAGQQAGQPNGSSQGKVHNPF
						LPTPMLPPPPPPPMARPVPLPVP
						DTKPPTTSTEGGAASPTSPTTRS
	į.	1		ŀ		/PGRTRPQQPFL/SYGPP*PHFLR
						AGIHHL
22037	52405	Α	22165	63	250	
22038	52406	Α	22166	1	817	SRLGHPRKDMDDDTKATLISY
1		1				MSLSFRHGYCFSEDPCVHTELG
		l				IQSGRGGRGDPDPPGNALE*LA
l						HLGP/SSHLGP*AQDGLVDAAG
1						RASPLAGTEGAGRFSHLQPI/GP
ł				ĺ		RGPPE*GAAGERFPSGVAG/GPR
1		l				GAHGLPCPEQLQLQPPPPS*MP
1		1	1	l	1	KTGSTFVAHFPSPLAQGHSPPG
				l		GGAGWCSGCSPDRPPLLGVGS
1						HGELLNPNSPAYEVGAG/RPAT
						GSSGTRAGPPSPSHSAPGRTPSV
			1			SLCD/RHQCLRASPTGSIGSLPA
1						HWREGLAGPLMD

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22039	52407	A	22167	240	1790	TAKNSYLESCPLIMENSIHQWP RQPFPPVRRQAADAACGAGSQ ALVPSHSDITDGVLPGAEWLGD GGPARVPGAAGGRGRATS*A GPLSL*APLCPSGLSGNGKQL LGAAGESSSHAQMHELHPAQ* HE*ATDHTGQGPPIG*KQNG PPCPDNDLPAPESTPPTWNEDF LPDAIPLAHPVPRQRRPTGPAGP GGCGPPGAPPGSKGDRGQT GEKGPAGPPGLLCPPGPRGLPG EMGRPPGPGPGPGPGPGRGPG SPCGALVSLQPPTDKDNGTQG WPLPSWTQCWSGVYGPRGPPG PGCPGPGPGPGPGFGSQGLA GERGTVGPSG/AGGAAAERGPE DPGRASPHPGAHDWDPRSPGLP RGFWPGCCPESOPQDEEGWH PITRUGPCCPSQPGDEEGWH PITRUGPCCPSQPCDEEGWH PTRWGPCCPSQPCDEGWH PTRWGPCCPSQPCDEGWH PTRWGPCCPSQPCDEGWH PTRWGPCCPSQPCDEGWH PTRWGPCCPSQPCDCAGAG PCGCAGAG P
22040	52408	A	22168	104	271	DKSQSAVSVLCARSRLRAQLRP GAEQMNDCKGITGTRQDIHAV SFASEHNIQNWVSKHLQLSPPV TPATQAPLKVRVPQASTLVL IPSCTNPYGGGRRCDSVVPNOCF
22040	52408	Å	22168	104	2/1	TKHMQSRLLG*GEKALEGGGG RGVKSLVTKRDF
22041	52409	A	22169	494	1074	IGTTLOCKHWGGGYLHWGUFE EMMRLTINIRSMDPKIMFAIWR VPSPFQIPTTRKSVGHRMGGGKI GAIPHYLTPVKAGRLUVEMG GRCEFARSAKVSLTOVALRSCP FASKRL*ACGTPREGCEQOQEE RERQOPRTPWTI*AE*PTAQHG WAYGKVPEPHMTLTPQGELG ARFYMPKTCVVSVGDILYIGY
22042	52410	A	22170	61	329	NQRSGKKSQPW*STITPPSLIWL SNSSTTLSSSKMGQSYSSS*PSY NCSVSRMFGSNMVPAATASGA SLTCPLLWLSMTHFRRKVGGW A
	52411	А	22171	21	444	STAQSISRSDLHYFLALLVYYW AADARFLPGRIPWLIHKLELEK HTEYCLVGSQYTPLVQVFRSN TLVSVLSTKTI*NQRSGKKSQP W*STITPPSLIWLSNSSTTLSSSK MGQSYSSS*PSYNCSSRAKSPTL TVFQS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hed	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide scauence	of peptide sequence	deletion, \=possible nucleotide insertion)
i				sequence		
22044	52412	Α	22172	163	1056	LEKSDHRISWSPGKTA\SPKSMP
		İ		l		ERWHR*WAQILKD\MGIT\EYEP
					1	R\VIN\OMLEF\AF\RYVTTIL\DD
						AKIYSSHAK\KATVD\ADDVPIG
						OSRC\RAGSSLFTSPPPKRF/LLL
				1		RYLAROKKSNPFCPLIKPY\SGP
						RLPPDRYC\LTAP\NYRLKSFTE
					ļ.	KGHSTSAGKELTVP\RLSVGFRL
				i		LSRTKVLPTLGAHPTPO\TMSVF
				ĺ		NLK*GTPHVPSPGQRFTSTRCPT
						S\OS\PAVKSFOFPGNLSOFONV
				l		L\INPIH*IGSQKHSYYPLNMMS
1						S\ONTA\NESSNALKRKREDDD
	ł			1		DG\DDDDDDDYDNL
22045	52413	A	22173	215	344	
22046	52414	Α	22174	302	1330	ACEQKRGGEGGNKNRERRKGE
	i					KAGRRQGTRQPCSRCMRRTKQ
1						VEKNDDDQKIEQDGIKPEDKA
		i		1		HKAATKIQASFRGHITRKKLKG
l					l .	EKKDDVQA\AEAEA\NKKDEAP
					l	/VFADGVEKKGEG\TTTARSSPQ
			ļ			PLAPKP\DEPGKAGETS\SEEKM
ŀ						GEG*LLATEQGSPPRLLAS\SEE
ŀ		l				KAGSAETESATKASTDNSPSSK
1						AEDAPAK\EEPKQADVPA/AFG
						VFPNPLGALLLAATTPAAEDAA
						AKATAQPPTETGESSQAEENIE
		1		ŀ		AVDE\TKPKESARQDEGKEEEP
						EA\DQEHAWNSGEMGFPHPHPP
						GCLACLSLPSSQLHSEVPSCPAH
ĺ	ł	1	l			VCESVLSHPLALFLSVWANI

SEO ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nuclcotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22047	52415	Α	22175	2	1577	HGFRLPHPQPALRRLLDQRRPK
						TEDRCGHWDQILDLSIIKHLFS
l						MSFIPVAEDSDFPIHNLPYGVFS
						TRGDPRPRIGVAIGDQILDLSIIK
						HLFTGPVLSKHQDVFNQPTLNS
İ						FMS\LGHAAWKEARVFLQNLL/
						SL*SQARLRDDTELRKCAFISQA
l				i		SATMHLPATIGDYTDFYSSRQH
		1				ATNVGIMFRDKENALMPNWLH
						LPVG\YHGRASSVVVSGTPIRRP
						MGQMKPDDSKPPVYGACKLLD
l						MELEMAFFVGPGNRLGEPIPISK
						AHEHIFGMVLMNDWSARDIQK
						WEYVPLGPFLGEEFWGPLSLP
						WGGCPWDAL\MPF\AVPNPKAV
						PQAPCRYLCHDEPYTFDINLSV
						N\LKG\EGM\SQAAT\ICKVQILK
						YMYWT\MLQQAHSPTLSNGC\N
						LRPGGPPGLSGTHQAGPEP/EKN
						FGSMVE\LSWKGTKPI\DLG\NG
						QT\RKFLLD\GDEVIITAMATRD
						FKWYC\QGDG\YRIGFGQ\CAGK
i						VLPALLPIMRFSLLFWKQRAQS
		_				TPFQPCDWGSSPRAVGLVPPFQ
22048	52416	Α	22176	3	623	DSRNSSILPRRGALLKVNQELA
						GYTGGDVSFIKEDFELQLNKQL
ł						IFDSVFSASFWGGMLVPIGDKP
						SSIADRFYLGGPTSIRGFSMHSI
						GPQSEGDYLGGEAYWAGGLHL
						YTPLPFRPGQGGFGELFRTHFFL
						NAGNLCNLNYGEGPKAHYIVK
						LAECIRWSYGAGIV\LRLGNIAR
						LIELNYCVPMGVQTGDRICDGV
			i			QFGAGIRFL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22049	52417	A	22177	352	2397	GPMSEGIASQGARRRRGIMGLR
22049	32417	^	22111	332	2391	GLWASRADGRRGETEPEAKQEI
		ı				LENKDVSFLLKVLIELLVETFTS
						IHH*GDDVFKAKNLIEVGVVST
ŀ						WCAFPVSSEELVNVPM*NIDTC
						QGTYCGVVSL*FPFCPSEKSCFY
						GQNNTMVGNNEGSMVCYRLF
	l					TFLYWNLELGTFTL*LHSKETS
						YGLSFFKPRPGNFERK*EAQQII
				l		EYTGLIEKLK*VAYL*KSAEYS
						VS\AFQSFPLGLGENRVAFSLIP
						KGSYLIQ*VSNGCWHLHCQPLL
						WGSYHRILAELRVLIESTWPFE
						G*LCNGTSQAFSCGPCTPSRSSA
İ	1	ı				RHGHSVCGQLLVADVPGVKG
				1		MLGENNHLFLKY*SMNG*FA/L
	İ					YFPLDLFSCFL*ESFPLTAPSLEG
						SRLTLETMYLLCQGGFGELFRT
	1					HFFLNAGNLCNLNYGKTCAIQE
				ŀ		TIVVQLFSCLKAQYTKIRLGNIA
						RLELNYCVPMGVQTGDRYVLG
						IIFHNHIPLSSNFILFWTWFASGD
						TAGAARLLRLPTDDRAFRDGG
1						GCSHNRQLPFPSTPLLFSTQSPS
1						WKGCSHAGCMHVCLGAQLPG
				l		LGAQLGHGSESQVERLLACGG
						RWGARRQGCCWHDLPKAPHA
1						GCFLAGEKKVGDDQEPCTKTA
						AVREGESVGCTAEGGCLGLSG
				1		GCQGWLAPGLVELRSAAARQE
				j		GSRRRRKRRRHLLRVHPQKVV
22050	52418	A	22178	1	1017	
22051	52419	A	22179	1	435	
22052	52420	Α	22180	1	1353	
22053	52421	Α	22181	1	834	
22054	52422	Α	22182	2	2675	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22055	52423	I A	22183	72	1339	AMRGANAWAPLCLLLAAATO
22033	32.423	<u> </u> ^	10J	1'-	1333	LSROOSPERPVFTCGGILTGESG
						FIGSEGFPGV\WPPESKCTWKIT
						VPEGKVVVLNFRFIDLESDNLC
		l				RYDFVDVYNGHANGORIGRFC
		l				GTFRPGALVSSGNKMMVQMIS
1		l				DANTAGNGFMAMFSAAEPNER
						GDOYCGGLLDRPSGSFKTPN\W
						PDR\DYPAGVTCVWHIVAPKNO
ŀ						LIELKFEKFDVERDNYCRYDYV
		l				AVFNGGEVNDARRIGKYCGDS
ŀ				ł		PPAPIVSERNELLIOFLSDLSLTA
1						DGFIGHYIFRPKKL\PTT\TEOPV\
ŀ				1		TTTFPVTTGLKPTVALCOOKCR
				1		RTGTLEGNYCSSDF\VLAGTVIT
						TITRDGSLHATVSIINIYKEGNL
						AIOOAGKNMSARLTVVCKOCP
						LLR\RGLNYIIMGPSRVKDGAR
						ONSWPNSFIMMFOT\RNOKLLG
						CP*KISQC
22056	52424	Α	22184	49	170	HTVGDGRLLRRP*REFGASFL*
						CFLEARFGLKNWGHCWRH
22057	52425	В	22185	597	651	
22058	52426	Α	22186	1	791	
22059	52427	Α	22187	2	943	TGACAEPRVGLLFRLKGRCRG
i						GRNMELGSCL\YGFREAAEEEG
l						EPEVKKRRLLCVEFASVASCDA
						AV\AQCFLAENDWEMERALNS
						YFEPPAEES\ALERRPETISEPKT
1						YVDLTNEETTDSTTSKISPSEDT
1						QQENG\SMFSL\ITWEYLMGL\D
ł						LKQSVQRRARGVCSYLAL*VLS
						LLFNSNVSYVYTERMNQLKMV
					ł .	LKKMQEAPESATVIFAGDTNLR
1						DREVSDKVSLMFEDRLLLWRLI
				1		*VPFCQYTWDTQMNSNLGITA
			l	1		ACKLRFDRIFFRAAAEEGHIIPR
1			l	l		SLDLLGLEKLDCGRFPSDHWGL
		_				LCNLDIIL
22060	52428	Α	22188	277	534	

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	sequence	or pepade sequence	detention, (-possible intelectine insertion)
22061	52429	Α	22189	29	1305	VGVVLRVECPLCPGPAPMSVPA
						FIDI\SEEDQAAELRAYLKSKGA\
						EISEENSEGGLHVDLAQIIEACD
						VCLKEDDKD\VESVMNSVGIPY
		l		l		SWIL\EPDKQ\EAFDLKALCEKA
1		i i				GSKFPRR*TAPSLETCSFLSNLF
		ı				H\GMG*GILPVKIHSVICSL\IKV\
1		l				AASCG\AIQYHPQLEPGIKFRK
						WDFLNWN\LTTGKKSTPLLKTT
		l				FMKALVE\CKKSDAASKGM\VE
		ı			ŀ	LLGSYP\EGNASQ\ARVDAHR\C
		l				IV\RALKDPN\AFLFD\HLLT\LKP
		l			1	VKFFGKASLFHCSFLTKF\VSA*
		1				MGHHMSRFYPD*LRTFH*IPLG
1		1				PVTMDPGIWPKMRLLF/TFMGN
1		1				GQ*KNKEISF*PQCQQEPSRLGA
		1]		D\DVESHLVIDA\VRN*KWVYC
		1		1		QNWSRPRRKVVVSHSTHR\TFG
						K\QQWQTTVLTHLNAW\KQNL
						NK\VKNSLLSLF
22062	52430	A	22190	194	413	
22063	52431	Α	22191	3	160	EVKFQWVLSLG*SFSLPKSSFQ
						RCGAGGAVHWSTALFLLGAAS
		_				ASSYHQRPG
22064	52432	A	22192	268	473	
22065	52433	C	22193	93	500 398	CEL DECL AT CODED COLONDAL
22066	52434	I ^A	22194	12	398	FFLRRSLALSPRPDCGLQWRNL GSLOAPLPGFTPFSCLSLPSSWD
		1				YRRPPPRPANFLYF**RRGFTLL
						ARMVSIS*PHDPPASASQSAGIT
						GVSHRARPETYNFCSFIIISSLPV
		ĺ				EPEARKMTKNSDIGEIW
22067	52435	A	22195	972	2180	PRHCCASASSAPAPSRPASSSSP
22007	32433	l^	22133	1772	2100	PAACPW*GSLLGPAATPATWR
						R
22068	52436	A	22196	2271	3074	RDPSTPTVVRSLGVPLLSRRRFK
		1				MAPSVAATKLAERNETMVOTC
						WTETQSSSDGSTPRRVRISLAL
		1		1		WRPSLFFSMADTSLVLNFPRVS
	l	l				SSGVIRPSNKRPSLAILTRISRTS
	1					SPKYMPLTIFSNSCFPGLRDPRSI
	1					SLSNSVRMSSNDPSSPKAPHS/G
ł		1		1		IDTHPPLVSFHOINVPOFLHVAS
1				1		IGASADDQADFTVLVVIATGHH
1	1			1	1	GAHCVIHHSYDVGIIVPSLFDGF
	1					VKQFDHICSFHSARFKSLCPGD
			ĺ	I		QDGLIYFGLLARKRKRESHR
						Y ODDING BUILDONN

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:		Nucleotide location of last Amino acid sequence (X=Unkn	
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	eodon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22069	52437	I A	22197	1956	2747	FAFSPK*HSCLRCPCI*FSSGLLH
22009	32437	l^	22177	1,750	2/4/	EVL*LLPLCWP*THGWDPGSRE
		1				ANKSPKLHAIRCVWVLEEENL
		1				WLSSNSOSLOTVKN*ESHIN*SC
	ŀ					RSNLI/HH*FWNOVK*K*LLNIS
	1					GNCFFFLRWSL/DSVAQAGVY
		1				WRDLGSLQPPPPGFKRFSCLSLP
		1				SSWDYWHLPPCLAKFCIFSRDG
	1				l .	GFTIWARLVLNS*SCDLPASGS
	1	ı				QSAGITGVSHHTW\LQVTYFLK
1						EMRSCYFSQVGWPQTPGLKQC
						SHLKLLSSWDYRHMSPHLAISG
22070	52438	c	22198	85	228	STILKELSS WD I KIIWISFIILAISO
22070	52438	A	22198	2	434	
22071	52440	A	22200	1	1356	
22072	52441	A	22201	146	1909	ILHSLEISALLSLMMLTIGDVIK
22013	32441	<u>۱</u> ^	22201	140	1909	OLIEAHEOGKDIDLNKVKTKTA
						AKYGLSAOPRLVDIIA\SVPPOY
1	1					RKVLMPKLKAEPIK\TA\SGIVV
1	1					VAVMCKPHRCPHISFTGNICVY
1	1					CP/GVGPDSDFEYSTQSYTG\YE
		1				PTSMRS/IFRARYDPFLOTKDRI
l						EO*ROLGHSVDKVEFIVMGGTF
	1					MALPEEYRDYFIRNLHDALSGH
						TSNNIYEAVKYSERSLTKCIGITI
		l				ETRPDYCMKRHLSDMLTYGCT
	1					RLEIGVQSVYEDVARDTNRGH
						TVKAVCESFHLAKDSGFKVVA
						HMMPDLPNVGLERDIEO\FTEFF
1		1				ENPAF\RPDGLK\IYPTLVIRGTG
	1					LYELWKS\GRYKSYFS**PGLN
1		1				WVARFLALVPPWTRVYRVQRD
		1				IPMPLVSSGVEHGNLRELALAR
	1	1				NEDLRIOCRDVRTREVGIOEIH
	1	1				HKVRPYQVELVRRDYVANGG
		1				WETFLSYEDPDODIL\IGLLR\LR
	1	1				KCSEETFRFELGGGVSIVRELH
		1				VYGSVVPVSSRDPTKFOHOGFG
	İ					MLL\MEEAERIS*RRNMGLGKI
	l					
		1				AVISGVGTR\NYY\GKIGYRLPR
			i			PRTMVKMLEIMGHTQSHSFCSI
22274	20110	١.	22202		400	LPWHEHRRIRISLNTQQRGL
22074	52442	A	22202	2	457 297	CUANDOVDI EL EMA OCUIONO
22075	52443	A	22203	l ²	291	CVMYRDYPLELFMAQCY/GNIS
		1		l	1	DLGKGRQMPVHYG/CKERHFV
1		1		l	1	TISSPLATQ/IPQAARGPGYGIM/
		1		l	1	SIRVDGNDVFAVYNATK/EARR
						RAVAENQPFLIEARPT

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 69/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \possible nucleotide insertion)
22076	52444	A	22204	3	1446	OWLLAKMAEAIAAARIVWRIN RGISHAALLLIRQLGGSGD*AR SHPPRQQQQCSALDDKPQPPG ASAGVIDKLEFIQPRVISGIPIYM VMDPQGGIIHQPQARDPHLAEG ERWLKLIVKSMTLL**HPWDRIL **VFSQGRILLIT**TNYWLRR GTHVOSARRLGQTBTIJVFGQY REJAGVLMYRDYPLELFMAQC VGNISHDLGKGRØMPWNYGCKE RHFVTISSPLATQDPSGRWGAA YAATGGANANRVR'VICYFRAR AGAS*GGDAHAGFINFRCHNL SCPHIFFCRNNGYAISTTTSEQYR GOGIAARGPGYGIMSIRVDGN DVFAVYNATKEARRRAVAENQ PPSHQAHDSRVQAPPSISDAS*T GYNYDENYWENKGPPILOF GHYLTSQGWWDEEQQKAWR KQSRKLPOPQYPSIVAEAEQ AERKVPKPNPINLLFSDVYQNM PAPVRQPPESLARHFKTYGNH YPWDHFDSK
22077	52445	Α	22205	2	292	

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SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \-possible nucleotide insertion)
22078	52446	A	22206		1894	QSLADLQSQPSITLTSQIWPSP AGFWSWGNLSVTPSKUPATP TRITAATPIQTILPQSQTAPKRID TPSLEEPSDLEELEQFAKTFKQR RIILGFTQGDAGLAMYKLYGN DFSPTTIFRFEALNLSFKNMCKL KPLLEKWLNDAENLSSDSS; SALNSPGIEGLSRRKKKTSIE TNIRVALEKSFLENQKPTSEEIT MIADQLNMEKEVIRVWFCNRR QKEKRINPPSSGGTSSSPIKAIFP SPSTSLVATTPSLVTSSAATTLS SPSLPLTSAAVTNLSVTASTSEA SSASETSTTQTTSTPLSSPLGTSQ VMYTASGLQTAAAAALQGAA QLPAMASLAMMAAAAGLNPSL MAPSQFAAGGALISLIPFGTLSG SLPTISLDATGNLVFAMAGGAP NIVTAPLFLIPQNLSLLTSNPPG GETIEISNNNSTMYMTGMRIQIG GTIEISNNNSTMYMTGMRIQIG GTIEISNNNSTMYMTGMRIQIG GTIEISNNNSTMYMTGMRIQIG GTIEISNNNSTMYMTGMRIQIG GTIEISNNNSTMYMTGMRIQIG GTIEISNNNSTMYMTGMRIQIG GTIEISNNNSTMYMTGMRIQIG GTIEISNNNSTMYMTGMRIQIG GTIEISNNNSTMYMTGMRIQIG TQAIERAPSYEIFGRTMQLNLS RSRWFOFFTREEALQADKKLN FYQCILLEQDSSCGTGPCSMTT YPYMLLSSEQPVQSLVRQCVI LLDAAAVISGGMGAVCLLLTH LCPLCYAGLLSEITDPESFVKDL

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SEQ ID			SEQ ID NO:		Nucleotide location of last	
NO:	of peptide	hod	in USSN 09/540,217	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22079	52447	A	22207	I	2417	MRVRTSSLPSTMLRVQNYARA
						KAGPEYAFVDSTECSPOWMVS
		l				PLKRGTOTNGLDFOKOPVPVG
						GAISTAQAQAFLGHLHQVQLA
						GTSLQAAAQSLNVQSKSNEES
		l				DSQQPSQPSQQPSVQAAIPQTQ
						LMLAGGQITGLTLTPAQQQLL
		l				QQAQAQAQLLAAAVQQHSAS
			l			QQHSAAGATISASAATPMTQII
			l			LSQPIQIAQDLQQLQQLQQQNI
						NLQQFVLVHPTTNLQPA\QFIIS
				,		QTPQGQQGLLQA\QNLLTQLPF
						QSQANLLQSQPRI\TLTSQPATP
						TCTIAATPIQTLPQSQSTPKRID
						PSLEEP\SDLEELEQFAKTFKQF
				1		RIKLGFTQGDVGLAMGKLYGN
			i	1		DFSQTTISRFEALNLSFKNMCK
			l			LKPLLEKWLNDAENLSSDSSL:
						SPSALNSPGIEGLSRRRKKRTSI
						EA\NIRVALEKSFLEN\QKPTSE
						ITMIADQLNMEKGVIRVWFCN
						RRQKEKRINPPSSGG\TSSSPIKA
						IFPSPTSLVATTPSPVTSSATTTI
		l				TVSPVLPLTSAAVTNLSVTGTS
	-	1				DTTSNNTATVISTAPPASSAVT
		1				PSLSPSPSASASTSEASSASE\TS
		1				TTQ\TTSTPLSSPLGTSQVMVTA
		١				SGLQTA/AQLLPFKGAAQLPAN
		1				ASLA\AMAAAAGLNPSLMAPS
		1				QFAAGGALLSLNPGTLSGALSI
		1				ALMSK\STLA\TIQTLASGGSLS
		_				K*TSLECKLGTLVICPMREEPPI
22080	52448	Α	22208	3	114	
22081	52449	A	22209	1	505	NLGVDLLPGYQDPYSGRTLTK
	Ì	1				GEVGCFLSHYSIWEEVVARGL,
		1				RVLVFEDDVRFESNFRGRLERI
		l				MEDVEAEKLSW\YSYWTLAYA
		1				LRLAGARKLLASQPLRRMLPV
		1				DEFLPIMFDQHPNEQYKAHFW
		1				RDLVAFSAQPLLAAPTHYAGD
		_				AEWLSDTETSSPWDDDSGR
22082	52450	Α	22210	3	626	RAARAAPLLHLLLLLGPWLEA
		l				AGVAESPLPAEVLAILARNAEF
	ŀ	l				SLPHYLGALERLDYPRARMAL
		l				WCATDHNVDNTTEMLQEWLA
		1	l			AEGDDYAAVVWRPEGEPRFYI
		l	l			DEEGPKHWTKERHQFLMELKO
		1	l			EALTFARNWGADYI/LGATLGI
		1		l		WQDHSVNSLGGCSPGRAGACI
	l	1	l			VPSSKFADTDNILTNNQTLRLL
						MGQGLPVVAPMLDSQTNV

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22083	52451	A	22211	I	2004	MRAARAAPLLQLLLLLGPWLE
						AAGVAESPLPAVVLAILARNAE
		l				HSLPHYLGALERLDYPRARMA
		1				LCLQGPAQPQWDLWLAGFEEE
		1				OTGTSEVHWGPTWOGSTTLEA
						SGHRCATDHNVDNTTEMLOE
		l				WLAAVGDDYAAAVWRPEGEP
		1			i	RFYPDEEGPKHWTKERHOFLM
						ELKQEALTFARNWGADYILFA
		1				DTDNILTNNQTLRLLMGQGLPE
						VAPMLDSQTYYSNFWCGITPO
						GYYRRTAEYFPTKNRORRGCL
						RVPMVHSTFLASLR\AEGADOL
		1				AFYPPHPNYTWPFDDIIVFAYA
						CQAAGVSVHVCNEHRYGYMN
						VPVKSHQG\LEDERVNFIH\LILE
						ALVDGPRMQASAHVTRPS\KRP
		[SKIGF\DEVFVISLG\RRPDRRER
		1				MLASLWEMEISGRVVDAVDG
		1				WMLQQQVPFRNLRP*DLLPGL/
1		l				YKDPYFGPHS*PRAEVGLLSSD
		1				HYSIW\EEVVSQGAWPR\VLVV
		1				*RNDVAPLRTNFQGGGLERL\M
	ŀ	1				E\DVEGKRNLSWEKPPFRILPLG
				1		R\KQVEPLRRET\AVEGLPG\LV
	1	1				VAG*LPNW\TLA\YALRLAGGP
		1				ASLLALQPFAPHCCPVERISLPI
		1				HCFDQ\HPNEASTKAHFWPRD\
	1	1				LVAFSAQ\PLLAAPTHYAG\DAE
	l	1				WILLSRRHHSPWDDDSGRLISW
		1				SGSQKTLRSPRLDLTGSSGHSL
22084	52452	Α	22212	1	468	ELSHNAKAKVMVASHNEDT/V
	İ	l				LFALRRMEELGLHPADHQVYF
		l				GQLLGMCDQI\SFPLGQAGYPIL
		l	1			TQIHLFSPHTCRAAGGEVRCLP
	1	I		l		ALPRVWALRCGPNLIPAWDSH
		l				WKLLGTLLELCAQGPHLCDPH
		l				VLGPRGLSTFCQGQPTQPEPLG
	1	I		i		EQWPGWG
22085	52453	Α	22213	1	1893	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1				sequence		
22086	52454	Α	22214	390	2005	INRDVETACTYILRNTEDVMLE
						FVMREWKKSRKLLGQRLFNKL
						MKMTFYGHFVAGEDQESIQPL
		l				LRHYRAFGVSAILDYGVEEDLS
	İ	1				PEEAEHKEMESCSSAAERDGSG
		l				TNKRDKQYQAHRAFGDRRNG
		ı				VISARTYFYANEAKCDSHMETF
		l				LRCIEASGRVSDDGFIAIKLTAL
		l				GRPQFLLQFSEVLAKWRCFFHQ
1			İ			MAVEQGQAGLAAMDTKLEVA
		l				VLQESVAKLGIASRAEIEDWFT
		l				AETLGVSGTMDLLDWSSLIDSR
		l				TKLSKHF/V*VPNAOTGOLEPLL
						SRFTEEEELQMTRMLQRMDVL
						AKKATEMGVRLMVDAEQTYF
		ĺ				QPAISRLTLEMQRKFNVEKPLIF
		l				NTYQCYLKDAHDNV\TLD\VDL
						SPREGWCFGAKPVRGAYLAQE
		ĺ				RACAAEIGYEDPINPMYEATNT
		i				MYHRCLDYVLEELKHNAKAK
		l				VMVASHNEDTVRFALRRMEEL
		ĺ				GLHPADHQVYFGQLLGMCDQI
						SFPLG\QAGYPVYKYVPYGPVM
		l				EVLPYLS\RRALE\NSSLMKGTH
						\RER\QLLWLELLRRLRTGNLFH
22087	52455	Α	22215	3	303	MERCIQUE II SEESIA CONTROLLER II
22088	52456	A	22216	798	2250	VPRAPOGML*GPRKDAGTPER
		l''		,,,,		RAGPPGGRGTADGGGCGALAG
	l					
	I	i				
			\$			SGS/DDPRLCIWTFLVTDQLLLP
						SGS/DDPRLCIWTFLVTDQLLLP TGPGRPRGSVAAAATGPGVGA
						SGS/DDPRLCIWTFLVTDQLLLP TGPGRPRGSVAAAATGPGVGA LRAPARAQPRGAVDGRPLLPQ
						SGS/DDPRLCIWTFLVTDQLLLP TGPGRPRGSVAAAATGPGVGA LRAPARAQPRGAVDGRPLLPQ VLPHRRLEVSLGKSPKECVLW/
						SGS/DDPRLCIWTFLVTDQLLLP TGPGRPRGSVAAAATGPGVGA LRAPARAQPRGAVDGRPLLPQ VLPHRRLEVSLGKSPKECVLW/ RHLLSESEKRPFVEEAERLRVQ
						SGS/DDPRLCIWTFLVTDQLLLP TGPGRPRGSVAAAATGPGVGA LRAPARAQPRGAVDGRPLPQ VLPHRRLEVSLGKSPKECVLW/ RHLLSESEKRPFVEEAERLRVQ HKKDHPDYKYQPRRRKSAKAG
						SGS/DDPRLCIWTFLVTDQLLLP TGPGRPRGSVAAATGPGVGA LRAPARAQPRGAVDGRPLLPQ VLPHRRLEVSLGKSPKECVLW/ RHLLSESEKRPFVEEAERLRVQ HKKDHPDYKYQPRRKSAKAG HSDSDSGAELGPHPGGAVYK
						SGS/DDPRLCIWTFLVTDQLLLP TGPGRPRGSVAAAATGPGVGA LRAPARAQPRGAVDGRPLLPQ VLPHRRLEVSLGKSPKECVLW/ RHLLSESEKRPFVEAERLRVQ HKKDHPDYKYQPRRKSAKAG HSDSDSGAELGPHPGGGAYYK AEAGLGDGHHHGDHTGQTHGF
						SGS/DDPRLCIW/TELV/TDQLLLP TGPGRPRGSVAAAATGPGVGA LRAPARAOPRGAVDGRPLLPQ VLPHRRLEVSLGKSPKECVLW/ RHLLSESEKRPFVEEAERLRVQ HKKDHPDYKYOPRRRKSAKAG HSDSDSGAELGPHPGGGAVYK AEAGLGDGHHHGDHTGQTHGF PTPLTTPKTELQQAGAKPELKL
						SGS/DDPRLCIWTFLVTDQLLLP TGPGRPRGSVAAAATGPGVGA LRAPARAGPGAVDGRPLLPQ VLPHRRLEVSLGKSPKECVLW/ RHLLSESEKRPFVEEAERLRVQ HKKDHPDYKYOPRRKSAKAG HSDSDSGAELGPHPGGGAVYK AEAGLODGHHHGDHTGQTHGG PTPLTTPKTELQQAGAFELKL EGRRPVDSGRQNIDFSNVDISEL
						SGS/DOPRICIWTELVTDQLLLP TGPGRPRGSVAAAATGPGVGA LRAPARAOPRGAVDGRPLLPQ VLPHRRLEVSLGKSRFECVLW RHLLSESEKRPFVEEAERLRVQ HKKDHPDYKYQPRRKSAKAG HSDSDSGAELGPHPGGAVYK AEAGLODGHHODHTGQTHGF PTPLTTPKTELQQAGAKPELKL EGRRPVDSGRQNIDFSNVDISEL SSEVMGTMDAFDVHEFDQYLP
						SGS/DDPRLCIWTELVTDQLLLP TGPGRPRGSVAAAATGPGVGA LRAPARAOPRGAVDGRPLLPQ VLPHRRLEVSLGKSPKECVLW: RHLLSESEKRPFVEEAERLRVQ HKKDHPDYKYOPRRRKSAKAG HSDSDSGAELGPHPGGGAVYK AEAGLGDGHHHGDHTGGTHGF PTPLTTPKTELQQAGAKPELKL EGRRPVDSGRONIDFSNVDISEL SSEVMGTMDAFDVHEFDQVLP LLGGPAPPEPGQAYGGAYFHAG
						SGS/DDPRLCIWTELVTDQLLLP TGPGRPRGSVAAAATGPGVGA LRAPARAQPRGAVDGRPLLPQ VLPHRRLEVSLGKSPKECVLW KHLLSESEKRPVEEAERLRVQ HKKDHPDYKYOPRRKSAKAG HKSDSDSGAELGPHPGGGAVYK AEAGLGDGHHHGDHTGQTHGF PTPLTTPKTELQQAGAKPELKL EGRRPVDSGRONIDFSNVDES SEEWMGTMDAFDVHEFDQYLP LGGPAPPEFGQAYGGAYFHAG ASPVWAHKSAPSASASPTETOP
						SGS/DOPRLCIWTEL VITOQLLLP TGPGRPRGSVAAAATGGVGGA LRAPARAGPRGAVDGRPLLPQ VLPHRRLEVSLGKSRFECVLW HKKDHPDYKYQPRRKSAKAG HSDSDSGAELGPHPGGAVYK AEAGLGDGHHHGDHTGQTHGF PTPLTTPKTELQQAGKPELKL EGRRPVDSGRQNIDFSNVDISEL SSEVMGTMDAFDVHEFDQYLP LGGPAPPEPGQAYGGAYFHAG ASPVWAHKSAPSASASPTETGP PRPHIKTEQPSGHYGDQPRGSF
						SGS/DDPRLCIWTELVTOQLLLP TGPGRPRGSVAAAATGFGVGA LRAPARAOPRGAVDGRPLLPQ VLPHRRLEVSLGSKPECVLW RHLLSESEKPPVEEAERLRVQ HKKDHPDYKYOPRRKSAKAG HSDSDSGAELDPIPGGGAV AEAGLGDGHHHGDHTGQTHGF PTPLTTRKTELQQAGAKPELKL SEEVMGTMDAFDVIBSEL SSEVMGTMDAFDVIBSEL SSEVMGTMDAFDVIBSEL GGAPPPEGQAVGGAYFHAG ASPVWAHKSAPSASASPTETGP PRPHIKTEOPRSGHYGOPPGGSP PRPHIKTEOPRSGHYGOPPGGSP DYGSCSGOGSSATPAPAPGFFAG
						SGS/DDPRLCIWTELVTDQLLLP TGPGRPRGSVAAAATGPGVGA LRAPARAQPRGAVDGRPLLPQ VLPHRRLEVSLGKSPKECVLW KRLLSESEKRPVEGEARELRVQ HKKDHPDYKYQPRRKSAKAG HKKDHPDYKYQPRRKSAKAG HKCHPDYKYQPRRKSAKAG FPTPLTTPKTELQQAGAKPELKL EGRPVDGGRONIDESNVDELSL SSEVMGTMDAFDVHEFDQVLP LGGPAPPEPGQAYGGAYFHAG ASPVWAHKSAPSASSPTFATED PRPHIKTEQPSGHYGDQPRGSS DYGSCSGQSSATPAAPAGPFAG SQGDYGDLQASSYYGAYPGYA
						SGS/DDPRLCIWTEL-VTDQLLLP TGPGRPRGSVAAAATGPGVGA LRAPARAQPRGAVDGRPLLPQ VLPHRRLEVSLGKSPFECVLW HKKDHPDYKYQPRRKSAKAG HSDSDSGAELGPHPGGAVYK AEAGLODGHHGDHTGGTHOF PTPLTTPKTELQQAGAKPELKL EGRRPVDSGRQNIDFSNVDISEL SSEVMGTMDAFDVHEFDQYLP LGGPAPPEPGQAYGGAYFHAG ASPVWAHKSAPSASASPTETGP PRPHIKTEQPSGHYGDQPRGSS DYGSCSGQSSATPAAPAGPFAG SQGDYGDLQASSYYGAYPGYA PGLYQYPCFHSPRRPYASPLIN
						SGS/DDPRLCIWTELVTDQLLLP TGPGRPRGSVAAAATGPGVGA LRAPARAQPRGAVDGRPLLPQ VLPHRRLEVSLGKSPKECVLW KHLLSESEKRPVEEAERLRVQ HKKDHPDYKYOPRRKSAKAG HKSDSDSGAELGPIPGGGAVYK AEAGLODGHHHGDHTGQTHGF PTPLTTPKTELQQAGAKPELKL EGRRPVDSGRONIDESNVDLSL SSEVMGTMDAFDVHEFDQYLP LGGPAPPEFGQAYGGAYFHAG ASPVWAHKSAPSASASPTETCP PRPHIKTEQPSGHYGDQPRGSS DYGSCSGGSSATPAAPAGPFAG SQGDYGDLQASSYYGAYPGYA PGLYQYCFHSPRRPYASPLLN
						SGS/DDPRLCIWTEL VITOQLLLP TGPGRPRGSVAAAATGPGVGA LRAPARAQPRGAVDGRPLLPQ VLPHRRLEVSLGKSRFECVLW HKKDHPDYKYQPRRKSAKAG HSDSDSGAELGPHPGGAVYK AEAGLODGHHODHTGQTHGF PTPLTTPKTELQQAGAKPELKL EGRRPVDSGRQNIDFSNVDISEL EGRRPVDSGRQNIDFSNVDISEL SSEVMGTMDAFDVHEFDQYLP LGGPAPPEPGQAYGGAYFHAG ASPVWAHKSAFSASASPTETOP PRPHIKTEQPSPGHYGDQRGSF DYGSCSGQSSATPAAPAGPFAG SQGDYGDLQASSYYGAYPGYA PGLYQYPCFHSPRRPYASPLLN GLALPPAHSPTSHWDQPRNSG TAQGPKEGVAGAQGAYRSKA
						SGS/DDPRLCIWTELVTDQLLLP TGPGRPRGSVAAAATGPGVGA LRAPARAQPRGAVDGRPLLPQ VLPHRRLEVSLGKSPKECVLW KHLLSESEKRPVEEAERLRVQ HKKDHPDYKYOPRRKSAKAG HKSDSDSGAELGPIPGGGAVYK AEAGLODGHHHGDHTGQTHGF PTPLTTPKTELQQAGAKPELKL EGRRPVDSGRONIDESNVDLSL SSEVMGTMDAFDVHEFDQYLP LGGPAPPEFGQAYGGAYFHAG ASPVWAHKSAPSASASPTETCP PRPHIKTEQPSGHYGDQPRGSS DYGSCSGGSSATPAAPAGPFAG SQGDYGDLQASSYYGAYPGYA PGLYQYCFHSPRRPYASPLLN

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
j	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
22089	52457	A	222.17	299	1631	I ILLEPHCGGKCFKRHWFPCSCEA
22009	32437	^	22217	299	1051	SC\PSAAPORW*HTG/SPEVPAA
1						PGAQGREPSM/RDGEGAVRGGS
1						CMEAGPMGLQCVDKKHQGVH
			İ			N/RGEARVFLOSOVGAILPRPW
1			l			PNPAPSEP/VQSPLASSTGP*VFP
1						SRPGOOASGPA/RRHPGAEPGO
ł			i		ŀ	RGDAVQLLPRPQTSPHGPQDA
1			ŀ			ALSSAPHPWRQL/MASPGLOEA
1	İ					
1						PLHPFCGHRSPCPVPWQLSPTD
						PIQSRPPARGWAPANQGRPSHP RCRGGGGAFCGHGF/SYTQPLT
			İ			
1						FRLASMVLSIHPKPPKETLRFRP
1						NMVVAAEPGVPAMSQRRPRGC
Į						TRFSHTDKAPLDTCTDMNAYT
						CARRHSSTYSDHITLQGAFACT
						HTWAQSCTCQPPPTPAAKCLDR
ŀ						GQSSGMLHAGCVGWGGGTHV
ŀ						RVDLSSQLNRRVRPHLCMLRC
						WSSCHRASVSCTHPSLHTTGPV
		<u> </u>		L		LISARTGADHTWYTAGTY
22090	52458	Α	22218	383	762	RHQGNQKRKELSLK/CQKSRQE
		l				SPRAEENPKWREGKKETSESSV
						QKAGRAAAAQAGAAASRVPG
		l				LSGSNLAPCNKGRLSAREDVSN
İ						SKLKILSSGRKHTEHVIMLKAW
						GLGRQSLRCRLFDLIPSHKV
22091	52459	Α	22219	325	945	QPSPLAPKADLSAREGRFQLKV
1	1		i		ľ	RCRLQQYQQQR\RKF\AAAFLA
						FHFSLLGSCGILAEAG\KKEKPE
l						KK\VKKSDCGRIGSWSVCVAPP
						SGDCGLGHHGRATRT*\AECKQ
						\TMKTQRCKIPCNWEGRQFGRG
				l		SAKLPSFQAWGECDLDT\ALKT\
						KTGSLKRALHKSPNAQKTVTIS/
				1		KSPCGKLTKPQNLQAES\KKKK
						KEGKKPEGRLRD

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22092	52460	Α	22220	1	1908	MKYIQDFQREKQEFERNLARFF
	1					EDHPDLIQNAKKSDIPEKPKTPQ
				1	ŀ	QLWYTHEKKVYLKVRPDATTK
						EVKDSLGKQWSQLSDKKRLK
į.		İ				WIHKALEQRKEYEEIMRDYIQK
					1	HPELNISEEGITKSTLTKAERQL
						KDKFDGRPTKPPPNSYSLYCAE
						LMANMKDVPSTERMVLCSQQ
					İ	WKLLSQKEKDAYHKKCDQKK
		1				KDYEVELLRFLESLPEEEQQRV
						LGEEKMLNINKKQATSPASKKF
						AQEGGKGGSEKPKRPVSAMFIF
						SEEKRRQLQEERPELSESELTRL
						LARMWNDLSEKKKAKYKARE
						AALKAQSERKPGGEREERGKLF
						ESPKRAEEIWQQSVIGDYLARF
			İ			KNDRVKALKAMEMTWNNME
				ł		KKEKLMWIKKAAEDOKRYERE
						LSEMRAPPAATNSSKKMKFOG
	1				ľ	EPKKPPMNGYQKFSQELLSNGE
					ļ	LNHLPLKERMVEIGSRWORISO
						SQKEHYKKLAEEQQKQYKVHL
	l					DLWVKSLSPQDRAAYKEYISN
	1					KRKSMTKLRGPNPKSSRTTLQS
						KSESEEDDEEDEDEDEDEEEE
	1				ŀ	DDENGDSSEDGGDSSESSSEDE
	٠.	i			ì	SEDGD/EGGADRAQWEGRGVL
						PGGPPGDSTSFLTPQNEEDDED
						EDDDEDDDEDEDNESEGSSSSS
		ĺ				SSSGDSSDSDSN
22093	52461	Α	22221	129	2365	
22094	52462	Α	22222	5	2730	PSLRFAISGFVWRLAAPASAAA
					i	ELRLDCSARRAPRCPRPALPSTR
		1				SAEGAVLCAFRREEQQAAATA
						AAAAAATAAAAAAPAPPPRP
		1				EEEPLPPREGAAAVPGRAGEGA
		1		i		ATQSQGGSRWSGKPGGRAAPA
		1		i		RPRRWLDSWRMNGEADCPTDL
		1				EMAAPKGQDRWSQEDMLTLLE
İ		1		1		CMKNNLPSNDSSKFKTTESHM
		1				DWEKVAFKDFSGDMCKLKWV
						EISNEVRKFRTLTELILDAQ*HF
	1	1		l		KNPYKG\KKLKKHPDFPK
22095	52463	Α	22223	1	446	RPSPPTSSM/PGLWGYSRPCSSG
		1		1		VPASALQA*SPAFSPASQRPPTT
		1		1		SWWEVP*EEGRA\PVWEGRIRA
						VGPAPAS*PSDLPAWSCRSWRE
		İ		1		TRG\SPAEDAMLLRAVEPLEGC
l		1	l	1		QRPLLDSEPP/SPARTSAPAASA
1	1	1		1		SPPAPGGPPCLARPLPR
			L		L	

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22096	52464	A	22224	582	1175	THTHVPAGTHPHTOTTSHSHLI
22070	32404	l^	ZZZZ-	302	1173	GRALTVPESPKCPCLLPSAC*PR
	1					PGRAOPAGPAWSKKEASNPGS
		ļ.				E/PPQHSL*RRMAPPE\PVPARA
	1					VHTCRCLSSAPPGARLPLDPER
		ŀ				RAPQSPRSPPPSAQPPPPPPPPPP
						PPPGRRGPSQQSWRPSPPPPRTR
	1	-				SRSPRRSRVRGPRAPALAAPPPP
	1					GARPGPCAAACGSPTGCS
22097	52465	A	22225	609	5241	GPOTTWCGRPTMCOHCGPAVS
22097	32463	I ^A	42223	1009	3241	SWKOPSHDPCCRGGENRROPG
		1				TPARATOGRLRRRPEGARVPSA
	İ					RRGRPRARWRDKERNNTCVA
ŀ		1				ML*QVTNSTAN\APQNNPEVDS
	Ì	1				IMAELKVEAPASVDWQKRCLT
		ı				LETQLFRFRLQASKIRELLADK
						GHGTYISPFSLAKQMQELEQRL
1						LEAEQRAENAETQVGVMEEKV
						KLSNLKNVDSEGSLHRKYQELL
		1				KAIKGKDELISQLEAQLEKQKQ
		L.			ļ	MRAEEAKTVQEKAAKIKEW
22098	52466	Α	22226	2	205	
22099	52467	Α	22227	I	6186	DI GOODING CONDUCTORIA
22100	52468	Α	22228	96	1584	DLGSGITASPPRPEPEHPVFPPA
					1	PHTPSKMGKEKTHINIVV\IGHV
					1	DSGKSTTTGHLIYKCGGIDKRT
						IEKFEKEAAEMGKGSFKYAWV
						LDKLKAEREPG\ITIDISLW\KFE\
						TTKYYITIID\APGHRDFIKNMIT
						G\TSQADCAVLIVAAGVGEFEA
		1				GISKNGQTREHALLAYTLGVK
		l				QLIVGVNKMDSTEPAYSEKRY
						DEIVKEVSAYIKKIGYNPATVPF
1	1	1				VPISGWHGANMLEPSPNMPWF
	1	1		ŀ		KGWKVERKEGNASGVSLLEA\
	1					LDTILPPTRPTDKPLRLPLQD\V
l		1				YKI\GGIGTVPVGRVETGILRPG
		ı		ł		MVVTFAPVNITTEVKSVEMHH
	l	1		1		EALSEALPGDNVGFNVKNVSV
1		1		1	1	KDIRRGNVCGDSKSDPPQEAAQ
	1	1		1	1	FTSQVIILNHPGQISAGYSPVIDC
	1	1		1	1	HTAHIACKFAELKEKIDRRSTK
1		1		1	1	KLEDNPKSLKSGDAAIVEMVPG
	1	1		1	I	KPMCVESFS\Q*PVIG/RTPVRD
	1			I	I	MRQTAA\VGVIKNVQKKSVGA
	1					GKVTKSAQKAQKAGK
22I01	52469	Α	22229	1	1023	
22102	52470	Α	22230	1	396	
22103	52471	Α	22231	2	1259	
22104	52472	Α	22232	1	2514	

SEQ ID			SEQ ID NO:			Amino acid sequence (X-Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	sequence	or peptide sequence	decenou, v-possible nucleotide insertion)
		i		·		
22105	52473	A	22233	3	2027	EESDCLTEYEEDAGPDCSRDEG
		l				GSPEGASPSTASEMDVSLHFVL
	1					WGCLHVYQRMIDKAEDVCLFV
		1				AQPGELVGQLAVLTGEPLIFTL
		l				RAQRDCTFLRISKSDFYEYDSP
						KLECGWHSEDPNPWTSPSTGL
	ŀ					NPLFLELCRHMVFQRLGQGDY
	ŀ	l			l	VFRPGQPDASIYVVQDGLLELC
		1				LPGPDGKECVVKEVVPGDSVN
						SLLSILDVITGHQHPQRTVSARA
					i	ARDSTVLRLPVEAFSAVFTKSA
	ľ					RTITGCVWRYVRASASYCPYLP
				l		PLCDPKDGHLLVDGCYVNNVP
						GSLWRYVRASMT\RRATCPRCA
	İ					TPRTGTYSWMAATSTICQASGR
						PHHPHTQAPPAPHTRTRRTSPA
				1		AWVPKRSSPLTWGARMRRTSA
	1				İ	PTGTACPAGGCCGSG*IPGLTR*
					1	RFQTWLKSSPAWPTCPVCGS*R
1						LSSPAPTASTCARPSTASRPWTL
1			İ			GSSTRSILRTTRRMSTAVTTTAT
	1					PAPIITPST*GTNGTGCAPQIRPA
						EPSPGARTSWNPSLSATFAPEEF
		1				VASP*LDVPICRMGQCCSLPGG
1		ł				ATLP*GPCGPQCMPGNVVPPER
						TAARYRQTTAYQTPACPTPPRG
				1		QRPT/CVPTVSSLQAGLQGGPRS
		1				DFDMAYERGRISVSLQEEASGG
1		1			l	SLAAPAREPREQPAGACEYSYC
1	1	1		l		EDESATGGCPFGPYQGRQTSSIF
1	1	1	I	l	i	EAAKQELAKLMRIE

2863

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion
	sequence		09/540,217	sequence	or pepnae sequence	acietion, v=possible nucleonae insertion
22106	52474	A	22234	281	1832	NDSGLPDRPSSWNQPMEAPLQ
		1				TGMVLGVMIGAGVAVVVTAV
		1				LILLVVRRLRVPKTPAPDGPRY
		l				RFRKRDKVLFYGRKIMRKVSC
	ŀ	l				TSSLVDTSVSATSRPRMRKKL
l l		l				MLNIAKKILRIQKETPTLQRKE
		ı				PPAVLEADLTEGDLANSHLPS
		l				VLYMLKNVRVLG\HFEKPLFL
		1				LCRHMVFQRLG/QGDYVFRPO
		l				QPDASIYVVQDGLLELCLPGP
		1				GKECVVKEVVPGDSVNSLLSI
	1	1				DVITGHQHPQRTVSARAARDS
	į.	1				VLRLPVEAFSAVFTKYPESLV
		1				VVQPGPPSGPAPYPAPSYGHA
		1				RALQIIMVRLQRVTFLALHNY
						GLTNELFSHEIQPLRLFPSPGLI
		1				RTSPVRGSKRMVSTSATDEPR
						TPGRPPDPTGAPLPGPTGLQG
		1				PRSDFDMAYERGRISVSLQEE
		1				SGGSLAAPAREPREQPAGACE
		1				SYCEDESATGGCPFGPYQGRO
				İ		SSIFEAAKQELAKLMRIEDPSL
		1	İ			NSRVLLHHAKAGTIIARQGDQ
		1				MEEVKVPAFAVIVD
22107	52475	Α	22235	50	4287	PGRKSPGAEDRALWGPACRL
		l				TDSGLPDRPSSWNQPMEAPLO
		1	Ì			GMVLGVMIGAGVAVVVTAV
		1	ŀ			LLVVRRLRVPKTPAPDGPRYR
		1				RKRDKVLFYGRKIMRKVSQS
			1			SLVDTSVSATSRPRMRKKLKN
		1	}			LNIAKKILRIQKETPTLQRKEP
						PAVLEADLTEGDLANSHLPSE
		1	ļ			LYMLKNVRVLGHFEKPLFLEI
						RHMVFQRLGQGDYVFRPGQP
		1		1		A\SIYVVQDGLLELCLPGPDGF
		1				CVVKEVVPGD\SVNS
22108	52476	A	22236	283	403	TCLSLVSGCGHCEKMNSEEN
				l		*IPIQWASSNPGRRRKNIR
22109	52477	Α	22237	33	222	NGPM\CFVKQLEIPQYGYRNN
		1				PTTTPRSNLAKELEKYSKTSFE
						TINDNHTYGGLGCMRPRLV
22110	52478	В	22238	1	1860	
22111	52479	Α	22239	303	672	GITACVFFSMAEGRTRAERSC
				1	I	SK*AIGAPAEE*GGASPGKSPE
		1				WQGAGYSPLGACGHKARLWI
		1				WRRSWSGEKGARGCRGTGK
		I				LGHQGWRWETPGRPGGAGPA
		1	1			GFPDPVVMGCEKEASRV
22112	52480	A	22240	I	1377	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22113	52481	A	22241		2232	MQDTEKDDNNNDEYDNYDEL VAKSILNLGKIAEDAAYRARTE SEMNSNTSNSLEDDSDKNENLG RKSELSLDLDSDVVRETVDSLK LLAQGHGVUSENMNDRNYA DSMSQQDSRNMNYVMLGRPM NNGLMEKMVEESDEEVCLSSL ECLRNQCFDLARKL SETNPOER NPOQNMNIKQHVRPEEDFFGRT PDRNYSDMLNLMRLEEQLSPRS RVFASCAKEDGCHERDDDTTS VNSDRSEEVFDMTKGNLTLLE KAIALETERAKAMREKMAMEA GRRDNMRSYEDQSPRQLPGED RRYSDSSDSHVKRPYYGKPPSRT EKKESKCPTPGCDGTGHVTGL YPHHRSLSGCPHKDRYPELLA MHESVLKCPTPGCTGGHVNS NRNSHRRDIQVFNLQKQEMIDR LESLSGCPHAAEKLAKQEKH MESVLKCPTFGCTGGGHVNS NRNSHRRDIQVFNLQKQEMIDR LESLSGCPHAAEKLAKQEKH LAKELEKYSKTSFEYNSONHT YGKRAIAPKVQTRDISPKGYDD ANGYCKDPSPSSSSTSYAPSSS SNLSGGGGSASSTCKSRSSPDYT HDMEAHMAATALNLSTRGR EMPQNLSTKPQDLCATRNPDGW EMPQNLSTKPQDLCATRNPDG EMPGNLSTKPQDLCATRNPDG EMPQNLSTKPQCUMPTYPTKRT EMPQNLSTKPQCMMNTRCF EMPQNLSTKPQCMMNTRCF EMPQNLSTKPQCMMNTRCF EMPQNLSTKPQCMMNTRCF EMPQNLSTKPQCMMNTRCF EMPQNLSTKPQCMMNTRCF EMPQNLSTKPQCMMNTRCF EMPQNLSTKPQCMMNTRCF EMPQNLSTKPQCMMTRCF EMPQNLSTKPQCMMTRCF EMPQNLSTKPQCMMTRCF EMPQNLSTKPQCMMTRCF EMPQNLSTKPQCMMTRCF EMPQNLSTKPQCMMTRCF EMPQNLSTKPQCMTMTRCF EMPQNLSTKPQCMTMTRCF EMPQNLSTK
22114	52482	A	22242	727	4372	\RIDEDESKDITPEDLDPFQEALE LWACVLPGVTLPPFLGEKTSCK MEVDTEEKRHRTRSKGVRVPV
						EPAIQELFSCPTPGCDGSGHVSG KYARHRSVYGCPLAKKRKTQD KQPQEPAKKRFAYVAN ASSSV DECDDSDGTEDMDEKEEDEGG EYSEDNDEPGDEDEEGDRE EEEELEEEDEDDEDGEDGEDVEDE EEEEEEEEDEDDEDGEDVEDDE EEEEEEEEEE

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22115	50400	-	22243	lad	1.000	DISCOURAGE HONOR DO LEGAL
22115	52483	Α	22243	34	1558	PASEEFNSLHGRTPRRDLECRN QQSTTEERRRHDCVQQASEHN
		1		}		EGKWDFAFSAQSQPKP*KPFDV
						PC*DSRPACP*EHTAGTSGSTSS
		ŀ				ETCFFQSSEWPYGELGAYNIFT
1						GPKSTLSSLVTGVSNSSSLS*LP
		l		•		SPKNTITNWFHTRDHSSYSEIHS
	İ					KACANP*APSQPSR*ASWIATG
		1		·		SFRRAGLAT*PCCTGSKLLPAW
	ŀ	l				VLANHRWDRTRDGIOKOGNSE
						*PSSPGTRAFEGNSSSPAPAASIT
						SMLSPSFTPTSVIRKMYESKEKS
						KEEPASGKAALGDSKEDTOKA
	İ					SEENLLSSSSVPSADRDYFSHY
						KFQTVRHYRGSSCS\PHLSQGQI
						FTPKEODYSDLKGNWGRKNTH
	ŀ					FGHSPVSLQHFFFRPVPPSSPLSF
						MVPNG*GLLTKLHPGFGTEDA
						GPGSTSTASSKFAPNWCASSWE
						GLESFTGNIWPHPGSALLPFN*S
	ŀ					LVHPLLNPRPGNTSASGNGATA
	i					ANSAQ\SCILQALVSHGSSCSAF
1	l					QDNPSETCPAGQACPHALP
22116	52484	A	22244	1502	2657	EGKVPLAFIVLTSLLNAVMSPSF
						FSCRLLVTALKVSSRCSAMKGV
		l				EFFTCWSTFKPFRPASTSSTEST
	ŀ					TPECELSLSFSLFAERLERRGFK
						STLAFCSISKISTLFSAWSSNGIG
		1				AK*FPEFCPGDRMACSRPASLSS
1		1				SSCG/S*PKTAGSAS*SARVREPF
						TEPTGRD*PFSQ\TSSIIEAKHGT
						LSRLKNSLKSKSPGDCSGKTAS
						LGTS*SAAGSCARMTSTSSSSAT
				1		PPLHSTIPSFTEAVRRLVLFLPL*
1				l		SSSSILSSKPVSSMVSD*LVGPA
	l			1		ENHSGSSSV*ESFLLRSPKTRLL
1	1			l		SPNSLLKRLSLKSLSRSLSLKSR
1		1		l		RSLSL\SGDPSQRSGQRLSSHCQ
1				l		SYVHQADEAHHYLSLMDFLSA
				l		GS*QRLSRGSPLQSSVSEG
22117	52485	Α	22245	324	391	
22118	52486	Α	22246	42	274	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
22119	52487	A	22247	1	835	MCAGNLCSWEERWAVDEKEA
		l				TDKKPHVGGFGQGAPAPAAPA
		l				AAMSGRSVRAVTRSRF*DDIKK
		l				VMAAIEKVRKWEKKWVTVGD
	ļ.	l				TSLRIFKWVPVTDSKEKEKSKS
		l			l	NSSAAREPNGFPSDASANSSLL
		l				LEFQDENS\NQS\SVSDVYQLKV
	ŀ	l				DSSTNSSPKPPKQSESLSPAHTS
		l				DFRTD\DSQPPTLGQGDPGRSPP
	i	l				LP\SSEVADEPPTLTKGRTSFHL
	İ	l			1	ETQVVEE\EEDSGAPPLKRFCV\
		l				DOPTVPOTGSEKLAPSRPSAFW
						ALPLFIAFWFWPRRVCWG
22120	52488	Α	22248	3	244	
22121	52489	В	22249	271	711	, and the second
22122	52490	В	22250	73	200	
22123	52491	Α	22251	885	975	GSTWWPAD*AQL*LHLPTRCM
						AECWEIPA
22124	52492	Α	22252	102	909	PGGRHGRQNDFAAPGGRLEARI
						HGTVSAPLYHRVLCRLLGSRPA
						SAAGPAARGGDLAPGEHSFCE
				1		DGNYARVESVDRSAINRAMQN
						PRKSTASVCWTFTASDLPAQGV
				l		\PRTSSTRTLREDPFPPSAEKWL
						RAVLHHFVNEKLQQIFIELTLKP
	l	l		l		SRRSMSGRHPLDSNPVLQQQGR
				l		CDLIENKLSPPGIMSVLDDVCPP
1	ł				l	CTPRAGEQTRHCCRTAGGCGD
1	I	1				PRAFQQLERRFVIHHYAGKTHR
	ł	l				PLPGLTPTGNSTNARNIVMDTQ
						QRTGNN
22125	52493	Α	22253	48	112	
22126	52494	Α	22254	1243	1615	YSPSSWFSNLYQPSTGALSSSET
		l		į.		PKDMAPMKSLREVRSQSHTSTL
ŀ				ł	I	SRRSTSPGFRLNTKVSFHTGFSV
1		1			I	AFTVRVFCFVRFRGSGMSFSFT*
					Į.	GSERPLGSIGIRLRASPTVTWISS
1	1		1	1	1	AVGARISSC

SEO ID	SEQ ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stup codon, /=possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1		l		sequence		
22127	10000	 -	1000.55	Lico	1431	la vagger a California de Livre de La California
22127	52495	A	22255	468	1431	SVSTTRSFSVDSSAKTAAMPVT
						VTRTTHHNHPDVIFGAWGPP*S
		1		l		WGPLWALTQPLGLL\RLLQL\V
		1				STCVAFSLVASVGAWTGSMGN
1	ŀ	1				WSMFTWCFCFSVTLIILIVELCG
	ŀ	1				APGPFPPWSWR\NFPI\TFACYA
		1				ALFCLSASIIYPTTYVQFLSHGR\
		1				SRDHAIAATFFSCIACGC\YATE
						VAWTRARPGEITGYMATVPGL
				l		LKVLETFVACIIFAFISDPNLYQ
						H\QPALEWCVAVYAICFILAAIA
		1				ILLNLGECTNVLPIPFPSFLSALA
						\FCLSSSMPPPLFSGPSTSSMRSM
	!	l				AASLGAREM*AAAAGMPTTW
		1				VPGTADWLRPS
22128	52496	Α	22256	98	414	CTRVSVAEKTSVCLFGLYWESL
i i		l				SISSSCESSAKWPFSTIRSASSTT
	İ	l				RHLSCRRFLK**CFSLINSHSRP
		l				GVATTISGRRFSILCCFCVDIPPT
						KIAVLIPVNLAT
22129	52497	Α	22257	2	224	
22130	52498	Α	22258	2	115	
22131	52499	Α	22259	2	118	
22132	52500	Α	22260	472	648	GPGGRSFTVWWNW*RRMAQC
						P\PAEGVQRPECLGPHRAVPAP
		1				AHVGLHQEPSRLLPALH
22133	52501	Α	22261	540	1893	GRCATPRPRWRGHQCDPPPGD
		1				VCVAWREMQLRNKKRRR\DLG
		1				NLIK\RFLTAPNSLPD/AVVLTTE
		1				EAVIFWCNVLAQQIFGL/PLEM
			İ			MNEQPLEGAVREKALHTMREQ
1		1				TQRMEGLVKQL/LTLSKIEAAPT
1		1				HLLNEKVDVPM/MLRVVEREA
1			İ			QTLVYNAVNHTPEGTHITVRW
						QRVPHGAEFSVEE*SEGWPVPV
		1				PKPHRAVVGANAHISFCPPQEL
1		l				SKRFTAIRKTKGDGNCFYRALG
						YSYLESLLPDAFOPLLPPOAWV
						SPGESQGAGIRDPLWPVPVPKP
		1		Į.		HRAVVGANAHISFCPPOELSKR
		1				FTAIRKTKGDGNCFYRALGYSY
		1	l	1	1	LESLLPDAFQPLLPPQAWVSPG
		1	l	1	1	ESQGAGTRDPLWLWLFRLWLL
		1		1	1	RLLCLLLLLGETAQSRLALTW
1		1	1		l	GRHKQGRLARLSGILGFLALRK
		1				QWLGQQVAQLLLQNLHLLLLR
						WAGLGVIVFQVSNQPAWERTG
22134	52502	A	22262	1	1772	Q.STITIQTSIQIAWERIO
22135	52502	A	22263	107	294	LHLLSFGYLTHQLLCLHLKWDS
-2.55		Ι''		1	I	HTLHHSHHTIAR*AGCSEATWS
1		1				PPRRLRAPRSRALCSGFR
		1			L	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22136	52504	A	22264	1	474	
22137	52505	Α	22265	1	148	VKYHEFTCSFKKQITVPEDARL LK*DVRGRPSRLFLADPPPPHT VNREY
22138	52506	A	22266	509	2755	YQSQSFDEDEMNQVTDWVDPS FDDFLECSGVSTITIATSLGVNN SSHRKNGPSTLIESSRFPARKIR GNLSSLEQIYGLENSKEYLSENE PWVDKYKPETQHELAVHKKKI IEIEVETWLKAOVLGROPKOGG
						SILLITGPPGCGKTTTLKILSKEH GIQVQEWINPVLPDFQKDDFKG MFNTESSFIMFPYQSQIAVFKE FLLRATKYNKLQMLGDDLRTD KKIILVEDLPNQFYRDSHTLHE VLRKYVRIGRCPLIFUSDSLSGD
						NNQRLLVPKEIQEEWSISNIRFN PWAPTIMMKFLNRIVTIEANKN GGKITVPDKTSLELLCQGCSGDI RSAINSLQFSSSKGENNLRPRKK GMSLKSDAVLSKSKRRKKPDR
						VFENQEVQAIGGKDVSLFLFRA LGKILYCKRASLTELDSPRLPSH LSEYERDTLLVEPEEVVEMSHM PGDLFNLYLHQNYIDFFMEIDDI
						VRASEFLSFADILSGDWNTRSL LREYSTSIATRGVMHSNKARGY AHCQGGGSSFRPLHKPQWFLIN KKYRENCLAAKALFPDFCLPAL CROTOLLPYLALLTIPMRKSSFR
						FSFYSKDIGEGLPSGRRTLWGRI GKWEGPGLDRGTWDD*TPDSG DEA\QLNG\GHSAEESLGEPTQA TVPETWSLPLSQNSASELPASQP OPFSAQGDMEENIII\EDYESDG
						DIEASLLIRLLHSFHFCFHFSGT
22139	52507	A	22267	2	161	
22140	52508	A	22268	I	599	VTRLFFQPGYGTLQPQRLLMPK KNRIAHYELLFKLEGVMVPKKI DVPIPKHPELA/DRNVPNLHVM KPMQSILKSIRGYVKLEQFA WIR HFYWYLTK*GYQGVSISSRDVL HLPPIEIVPCHPYGSPSRRLAR PSG*KVLEG*ANLRGLHKNGEA DKRLPTRRECLCHPGA/DKKAE AG/ALGSNPNFQFRGGGRGR GOJPPQ

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first eodon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22141	52509	A	22269	1	345	KHPELADKNVPNLHVMKAMQ SLKSRGYVKE\DYLHLPPEIVPA TLRRSRPETG/RPRFKGLEGERP ARLTRGEADRDTYRRSAVPPG ADKKAEAGAGSATEFQFRGGF GRGRGQPPQ
22142	52510	A	22270	2	161	
22143	52511	Α	22271	2	641	RRPQHRACSFFFSPGTGFCSRR CLMPKKNRIVAIVELLERGEGVM VAKKDVHMPKHPELLADIKNG APTFHVMKGHAGLSKSPRATV KGTSFAWGDIFYWYLTN-GY QGVSISSRDYLSSCPREICCLAT ATARSPSKRIGKAFGLKRSV GLRRPGEDFTRGGKFDRGVLT GRIAVATLVADKKAEG/ALGS SPPINFQFRGGFGRGRG/PPQ
22144	52512	A	22272		1693	MINSYPYQNEOGSNFEGYKA EFRPOTOTOEVIKGMEDSAFIFEGYKA EFRPOTOTOEVIKGMEDSAFIFED TVNREVTTDNPYTISVTNKTLS ARIKIMFMPRAYELKYTVIKM AFELSTRESKERM/DGSFETVLT DVIEGKTMSGYDRSRVNLPNF NNQVIFRVYRKTPDSNDSNVV DAIQVRSYAEVIDAKFRYPLTG LLFVEFDSKMFPNQLPTISIRKR WKINNYPSNYPDESRTYNGNW DGTFKKAWTNNPAWVLYDLM INQRYGLDQKELGISVDKWAL TYEYACAGYCDOMVULTAKAARNL ATEVT/CIIQSQTDAPREPAYIET RAT*CLMMSKTCINRTLSQYSI/ V/EGHSTVRNNVTSITAIGAHTV EA/KANRIGRWILKTNLRSHTV PATGLEGMIPTIGDVAISDNF WSNLTMNLSGRLLEVSGSQIF LFFRVDARAGDFIIVNRPDGKP VKRTISSVSADGKTEINIGFGFP VKRTIFABATGTR KONTONIONIONIONIONIONIONIONIONIONIONIONIONIO

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22145	52513	A	22273	I	1843	MEDNLISINKIKILLSVSDGEIDE
						TFSLKQLMFNSVPVQNEDGSFN
	ļ					FEGVKAEFRPGTQTQEYIKGME
						DSSSEVTVNREVTTDNPYTISVT
	l					NKTLSAIRIKMFMPR\AYELKVT
						VIKMAFELSTRCSKRLM/DGSFE
						TVLTDVIEGKTMSGYDRSRRV
					İ	NLPNFNNQVIFRVVRKTPDSND
						SNVVDAIQVRSYAEVIDAKFRY
						PLTVLLFVEFDSKMFPNQLPTIS
						IRKRWKIVNVPSNYDPESRTYN
					}	GNWDGTFKKAWTNNPAWVLY
						DLMINQRYGLDQKELGISVDK
						WALYEAAQYCDQMV\LTAKA
						ARNLATFVT/CIIQSQTDAYKVV
						RDICSIFRGMSFWNGPREPAYIF
						TNDNVVNGDFSYT\SQARRACT
						RRAT*CLMMSKTCINRTLSQYS
		1				I/V/EGHSTVR\NNVTSITAIG\AH
						VEA/KANRRGRWILKTNLRSTT
						VNFATGLEGMIPTIGDVVAISD
						NFWSSNLTMNLSGRLLEVSGSQ
						IFLPFRVDARAGDFIIVNKPDGK
						PVKRTISSVSADGKTIEINIGFGF
						PVKPNTVFAIDRTDLALQQYVV
		1				TKIDKGDDDEEFT\SKSRRWSTI
						LISTMKLITGLTSTTDRQASLSQ
						IR/CPKPENVK\CPQSRESSRG
22146	52514	В	22274	1	1452	

2871

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	l			sequence		
22147	52515	A	22275	I	1902	MIOKVISGSKGGSQKPHNPVEM
						EDNLISFNKIKILLAVSDGEIDET
		l				FSLKQLMFNSVPVQNEDGSFNF
		1				EGVKAEFRPGTOTOEYIKGME
						DSSSEVTVNREVTTDNPYTISVT
						NKTLSAIRIKMFMPR\AYELKVT
1						VIKMAFELSTRCSKRLM/DGSFE
l						TVLTDVIEGKTMSGYDRSRRV
		1				NLPNFNNOVIFRVVRKTPDSND
			1			SNVVDAIQVRSYAEVIDAKFRY
			ŀ			PLTGLLFVEFDSKMFPNOLPTIS
						IRKRWKNVNAPSNYDPESRTY
		l			ì	NGNWDGTFKKAWTNNPAWVL
						YDLMINORYGLDOKELGFSVD
						KWALYEAAQYCDQMV\LTAK
						AARNLATFVT/CIIQSQTDAPRE
						PAYIFTNDNVVNGDFSYT\SOA
						RRACTRRAT*CLMMSKTCINRT
						LSOYSI/V/EGHSTVR\NNVTSIT
						AIG\AHVEA/KANRRGRWILKT
			İ			NLRSTTVNFATGLEGMIPTIGD
						VVAISDNEWSSNLTMNLSGRLL
		1				EVSGSOIFLPFRVDARAGDFIIV
		1	İ	i		NKPDGKPVKRTISSVSADGKTI
						EINIGFGFPVKPNTVFAIDRTDL
		l			1	ALQQYVVTKIDKGDDDEEFT\S
		1				KSRRWSTILISTMKLITGLTSTT
		ı	Į.		i	DROASEVKPVNFTSSRHOAOPE
		1				VGLIPPPLWPQGKGIIAQTSKSS
22148	52516	A	22276	1	675	Total Carrolling Carroll
22149	52517	A	22277	155	700	SGVLVGHALTAROPSSGRRLV
		ľ				VDHSANRRGRWILKTNLCSTT
		l				VNFATGLEGMIPTIGD/GDFIIVN
		l				KPDGKPVKRTISSVSADGKTIEI
		I				NIGFGFPVKPNTVFAIDRTDLAL
		l				QQYVVTKIDKGDDDEEFT\SKS
1		I		l		RRWSTILISTMKLITGLTSTTDR
		l				QASLSQIR/CPKPENVK\CPQSRE
		I				SSRG

NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22150	52518	Α		1	1871	MEDNLISINKIKILLAVSDGEIDE FTSLKQLMFNSVPVQNEDGSSN FEGVKAEFRØTGTGTGEYIKGME DSSSEVTVNREVTTONPYTISVT NKTLSAIRIKMFMFRIAYELKYT VIKMAFELSTRCSKRLM/DGSFE TULTDVIEGKTMSGYDRSRRV NLPNFNNQVIFRVVBKTPDSND SNVVDAIQVRSYAEVIDAKFRY PLTGLLFVEFDSKMFPNQLPTISI IRKRWKIVNYPSNYDPESRTYN GNWDGTFKKAWTNNPAWVLY DLMINGRYGLDQKELGISVDK WALYEAAQYCDQMVLTAKA ARNLATFVTCIIQSQTDAYKVV RDICSIFRGMSFWNGESISVIID RTEPRAVITIONVVNGPSYTY SQARRACTRRAT*CLMMSKTGI RTLSQYSIV/EGISTVNINNT SITAIGAHVEARPTDCGRWILK TNLRSTTVNFATGLEGMIPTIGD VALSIDNEWSNLTMNLSGRLQ EVSGSQIFLPFRVDARAGDFNIY KPDARFVKRTISSVSADGKTI ENIGFGFPVKPNTVFAIDRTDL ALQQYVVTKIDKGDDEEFTIS KSRRWSTILISMILGTSTT DRQASLSQIR/CPKPPENVKCPQS RESSRG

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SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22151	52519	A	22279	3307	5305	CRSSONTQETSPDRALLVNLRP SQOPLLALWYPAEGVGGKGHG LPPRSFFRMDNQLVPAVSDGEI DETSLKQLMFNSVPVQNEDGS FNEGVAKERPOTTOTGEVIKG MEDSSSEVTVNREVTTDNPYTI SVTNKTLSAIRIKMFMRNAYEL KVTVIKMAFELSTRCSKRLM/D GSFETVLTDVIEGKTMSGYDRS RRVNLPNRONQVIFRVYRKTPD SNDSNVVDAIQVRSYAEVIDAK FRYPLTGLLFVEFDSKMFPNQL FNISIKR WKINVPSNYDPESS TYMGNWDGTFKKAWTNNPAW LYDLMINQRYGLDQKELGISV ULYDLMINQRYGLDQKELGISV ULYDLMINQRYGLDQKELGISV VXDLCSIFRGMSFWNGESISVII DRREPAYIFTNDNVVNGDFSY TSQARRACTRRAT*CLMMSKT CINRTLSQYSIV/EGHSTVRINN VISITAIGATYRIN VISITAIGATYRING VI
22152	52520	Α	22280	3	412	PAAVA*VKTEPETPGPSCLSQE GQTAIVKTEESSELGNYVIKIDH LETIQQLLTAVVKKIPLITAK\W QRAMTMRKVLQEILEKNPRFH HLTPLKTKHIAHWCRCHGYTPP DPESLRNDGDSIEDVLTQIDSEP
22153	52521	A	22281	2	513	IFACWGKQAWPACCNSLRGQA VKGNQLLPVSLVKRKTTLAPNT QTASPRALADSLMQLARQVSR LEVGRAKREPENEEEVDILSLSE PVKINIKKEQEEKQEEVKFYLPP IFGSEFIGDVTQKRLEEVDSTFG DDLVQGFSGGISRRYGEKPGVQ DLEVEPDSVTEMLQFYD DLEVEPDSVTEMLQFYD

SEQ ID NO:	of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22154	52522	A	22282	1	1197	MAWPCISRLCCLARRWNQLDR SDVAYPLTLHGYSDLDSEEPGT GGAASRRGQPPAGARDSGRDV PLTQYQRDFGLWTTPAGPKDPP PGRGFGAGGRGKSSAQSSAPP PAGARGVYLPIGDADAAAAV TTSYRYGLGRDRNESPPCGGRT RRPGARGMGWAAERRLEPGS HATSELPASEVTPWSVGTA GGAFAAPCPETVLEHPRAGSAP LPSQPPSWGQPSEWFAFSRVGT GLPLTPTAGSRARGARPCP ALPGHCLLDRTYTGLOTLGAET LLAVVNSAAMNVGVQVVDVE LIKHSLGEDCIYPQSSESDISDA
						PPSLPLTIPAPVKASSPIKQSHEP VPDTSVEKGS\PGSCPFHL*GPL SHLGSSPGFLLWRPPGLLSSVA
22155	52523	A	22283	131	677	PSSLS/CDIFLRSPISTPSPSPLPRT TYSTPYHVKQGTAGSVINPPYV IMDKQPGQVIGATTPSTGSPTN KISTASQISQGTGSPVPKIHGSSF VTSTVKVIIKQEPGEAPHVPAT GAASQSPLPQYVTVKGGHMIA VSPQKQVITPGEGIAQSAKVQP SKVJCQGIG*CLPTLARADLLYS
22156	52524	A	22284	3	5128	LLKHISSQSSPQAAVLTIPSQLK PLSYNTSGGVQTILMPVNKVVQ SFSTSKPPAILPVAAPTPVVPSS APAAVAKGEDASCFSAKSVEQ YYGWNIGKRRAAEWQRAMTM RKVLQEILEKNPFFHHLTPLKT KHIAHWCRCHGYTPPDFESLEM DGOSIEGVLTQIDSEPDHGSLG DGKSSSVGPTAVVTLLACHLEC PSSFSSADNLCRKLEDLQQFQK REPENEEEVDILSLSEPVKINIKK EQEEKQEEVKFY
22157	52525	Α	22285	112	651	GGLCTMARTKADWPANRTGG KAPIRKQLAYKKPASQECGPLT GGYKKPHRYRIPGTVGGSVKFR RIYQKSTIELLIRKLPFG/RRI.VR ELAQGL*NKILRFQSAAIGIALOG GKWRPHLVWPFLKDTNI.VC*S MARKYTIIAKKTFQLAPPHTVE NRALRIHSWMGKHFISQKKKK NFSSSCYW
22158	52526	A	22286	1	339	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22159	52527	Α	22287	2	1369	EPAPPARRSLSAAAAASGTTVD
	1	l				TASHSLVFSQQ\LNMQREIGFRV
		1				NCTVAM*DVYFKAHRAVLAAF
	1		1	ł		SNYFKMIFIHQTSECIKIQPTDIQ
	1	1				PDIFSYLLHIMYTGKGPKQIVD
	İ					HSRLEEGIRFLHADYLSHIATE
	l	1		!		MNQVFSPETVQSSNLYGIQISTT
	1	1				QKTVVKQGLEVKEAPSSNSGN
	İ	1				RAAVQGDLPQLQLSLAIGLDDG
	1	1				TADQQRACPATQALEEHQKPP
	1	ł		ĺ	i	VSIKQERCDPESVISQSHPSPSSE
		1			l	VTGPTFTENSVKIHLCHYCGER
	ŀ					FDSRSNLRQHLHTHVSGSLPFG
ĺ	ĺ		İ			VPASILESNDLGEVHPLNENSE
						ALECRRLSSFIVKENEQQPDHT
		ļ				NRGTTEPLQISQVSLISKDTEPV
						ELNCNFSFSRKRKMSCTICGHK
						FPRKSQLLEHMYTHKGKSYRY
		1				NRCQRFGNALAQRFQPYCDSW
		l				SDVSLKSSRLSQEHLDLPCALES
						ELTQENVDTILVE
22160	52528	Α	22288	363	947	STARRGKSRSRPAVALQRQPSA
		l				ASRSPSSSRRRRGVSKEVSVPLA
		l		1		RTKQTARKSTN\GK\APRKQLA
		l			i	YKKPARKSA\PSTGGVKKPHRY
				}		RPGYCGRSVKFRR\YQKSH*DF*
		l		Ī		FRKLPFQAVWVRENLLKDF*K
						QILRF\QSRSYRVALQGGQVEA
						LSGVGLF*KTTNLLC*SMPKRV
		1				NNYCQKDIPAKHARIPWRTVL
22161	52529	Α	22289	1	92	
22162	52530	С	22290	10	198	
22163	52531	В	22291	74	205	

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SEQ ID SEQ ID NO: Met SEQ ID NO: Meteodide No: well-optical to lead Amino acid sequence No: optical to lead No: optic	ble nucleotide
	cleotide insertion)
22164 52532 A 22292 2 1914 RGHCEPSRAAP:	
ARDPGRRRGPA	
	GQAAPQPMEP
ENIVANTVLLK	AREGGGGNRK
GKSKKWRQML	QFPHISQCEEL
RLSLERDYHSLO	CERQPIGRL\LF*
EFCATRPELSRO	IAFQDGMAEY
EATPDEKRKAC	GRRLMONFL\S
HTGPDLIPEVP\	
EQGPCKELSQE	TRVTHEYLSV
APFADYLDSIYE	
REAI\TKNTLRO	
EVCACOVRATO	
KKRIKKRKGEA	
KVN\SRFVVSLA	
CLVLTLMNGGE	
AGFPEARAVFY	
HRERIVYRDLK	
RISDLG\LAVHV	
\TVGYMAPECL)	
WALGCLLYEM	
KKIKRE\EVERL	
FSPQARSLCSQL	
CRGGSAREVKE	
LGAGMLEPPF\K	PDPQAIYCKD
VLDIEQFSTVKC	VELEPTDQTF
YQKFAT\GSVPI	PGKNEQMVET
ECFQELNVFGLI	OGSVPPDLDW
KGQPPAPPKKG	LLQRLFSSPKD
	LPYPPLAPSPRP
22165 52533 A 22293 102 403	
22166 52534 A 22294 1086 1436	
22167 52535 A 22295 965 1231	
22168 52536 A 22296 1434 2215 TPSFSLRTGVNK	EHWVLLGEST
EDGRW\MFTRG	
LRSRNLOCORIF	
LHPN\QAELIVG	
KTDHNEQLIPER	
DASLHGQLFNS	
GG\IGDEVTQFII	
ALQCRFSPDSTI	
CKIWRTSNFSLM	
GE\SSRG\WMW	
VTASSNNLARL	
YGGHQKAVVC	PFNDSVLG
22169 52537 B 22297 158 385	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nuclcotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
22170	52538	A	22298	145	1232	LPVRAEPTRAAAMSGDEMIFDP
						TMSK/NMKKKKKKPFMLDEEG
					i	DTQTEETQPSETKEVEPEPTED
						KDLEADEEDTRKKDASDDLDD
			1			LNFFNQKKKKKKK\TKKIFDIDEA
						\EKGVKDL\KI*SDVQEPTEPED
				İ		DLDIMLGHKKKKKKKNVKFPDE [*]
				1		DEILEKDEALEDEDNKKDDGIS
	ŀ	1				FSNQTGPAWAG\SERDYTYEEL
						LNR\VFNIMREK\NPGYWLPGG
						RKRKFGPWKPPQV\VRVGTKKT
				İ	i	FFCPLLQDICKLLHRSAQSISFA
						FLLG*IGVPSGSIDG**PNL*SKG
						RF\QQKQIENVLRRYI\KEYVTC\
				1		HTC\RSPDT\IL\QKDTRLLFPYS
		1				AETC\HS\RCFCWPVFKTGLPQA
						VHGQSRGTAPVPKA
22171	52539	В	22299	50	319	
22172	52540	A	22300	3	235	LLLYPLRSLPENPLRLVQQIYSF
		1	1			LKGSSEK*NDLPRVPLLELGVG
		l	1			LLPPNQYSCSRAKPCLTCGLTM
		_				SLDGAQQGFW
22173	52541	A	22301	3	2849	WAKPVARASPAAVPGRQCQIR
		l				GAWSWGEGRGQPGPAPSPAGS
		1			1	PATSEESLAPMAWRCPRMGRV
	1	l				PLAWCLALCGWACMAPKGTQ\
		l			İ	AEESPFVGNP\GNITGARGLTGT
		l				LRCQLQVQGEPPEVHWLRDGQ
		l			İ	ILELADSTQTQVPLGEDEQDDW
						IVVSQLRITSLQLSDTGQYQCLV
1		l		l	ŀ	FLGHQTFVSQPGYVGLEGLPYF
						LEEPEDRTVAANTPFNLSCQAQ
				I		GPPEPVDLLWLQDAVPLATAP
20174	50540	١.	22202		3899	GHGPQRSLHVPGLNKTS LVFLFDDLGIVELVLPGNIVHGI
22174	52542	A	22302	3364	3899	
		1				MSVRPRVAAGLRVRLQPGRRW
		1				LRGPLQLGAPLGLDLLVDVPGS
		1				PRFVVGVELPLVV/EAPRPPTPP
						AAAQQARS*GLECPGRGAAAP
		l		1		REVLHLQTAHIHGPGPRGGPR
		l	1			ARPGLPGLCPP*T/FSCQEAKGE
		L		L	L	RGGAHSNGGSGSKWPPPPPPPS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stnp codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22175	52543	A	22303	1660	4652	PNPNTCLCNHVESGPCCPSTHT
						YTLTDLOPLFGVRVPTRPSGRG
	1					TOAEESPFVGNPGNITGARGLT
		l				GTLRCQLQVQGEPPEVHWLRD
		1				GOILELADSTOTOVPLGEDEOD
		1			1	DWIVVSQLRITSLQLSDTGQYQ
	1	ı				CLVFLGHQTFVSQPGYVGLEGL
	i i	l		l	l	PYFLEEPEDRTVAANTPFNLSC
						OAOGPPEPVDLLWLODAVPLA
	1					TAPGHGPORSLHVPGAAVGAW
	1	l				EGGSRLWLRAPEAGRKRWGGE
		1				VPPVTHAGAGMLRSERD
22176	52544	A	22304	54	288	SEAGQAQPAAAVGFVWTGRRA
		1				WETGEGYYSNSLSMELQL*RQ
	1	1				CI*DCFFSSSWRCSVPVYFKEEK
	i .	1				YKGNLVCLLFFK
22177	52545	В	22305	243	2641	
22178	52546	Α	22306	35	204	
22179	52547	Α	22307	244	424	PSLFKNIMTFYPNLHFLWCRGR
		l				LYLDVWRQ/VR*FVRNTNYOLP
		l				LVLHNLLSRFNTFDELV
22180	52548	Α	22308	1	477	
22181	52549	Α	22309	2	418	WYQHLLLMRASGSFQSWWKV
		1		1		KGEPAYHMANRP**SLISLRTK
	1	l				AKTGA*LFDYFVPHFIRKN*SLF
		l				TIAKAWNHPKCTSVTDWIKKM
				i	İ	WYIYTMEYYAVTRRNKIVSFA
	1	ĺ				ET*MELEVIIVSKLTQEQKTKHC
						MFSLTSGS
22182	52550	Α	22310	30	199	
22183	52551	Α	22311	547	682	LVFLTNQRLVSTHQGTW\WE*K
						*CQPVIPSL*FYFLRKFSVIVIW*
		1				*LLILP*HPMALKI/C/CCQAACS
i		l				GHFTAGTGEITRPCSISSKVLKR
		1		l		LSRL*STSGS**LVFLTNQRLVS
		ł				THQGTWMKLEAIILSKVTQEQK
				l		TKHHMLSLRSGS
22184	52552	Α	22312	2	194	CPSMIDWLKKMWHRYTMEYY
		l				AA\IKKNEIMS\FAGTWMKLEAI
						ILSKLTQEQKN\KHCMFSLITGS
22185	52553	С	22313	1	1845	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				scquence		
22104	I so s s s	Ļ	00014	1	1384	L APPODRTUOUS VINCES COLUMN
22186	52554	Α	22314	1	1384	MRFQDRTWQVPYPSQCSWGSS
						RQGYVPQARGIVLDSQEAYPSP
		1		l		GGTTLCMHDPDKQAPG/PVRRP
				l		APGN\PNRLVVSVALQFLTFHES
						GITSALRVRPLPQSV\SFQGPGC
		l				TEA*RRASICRRRVS*HSTKLDL
		1				GAVAAQQQAWVAG\PRSPQRR
		ı		l		CHQYS*AGPAWLGLQRGRGSH
		l			l	RPDTSPWHWFWALAGREEQVL
		l		l		PPESLL/RARTPGTHLAPGPGA*
		1		l	1	TRVQSGPGSWLTWAG*GTPQVI
		l		1		SS/PGAGAVGAPETTAVPTV\$IA
		l		i		PRRSVEMWPETKEHLVAVARR
		1				KWVSAQEKRLSLRGCGSEKMG
		l		l		VSPGETRVYFPALTS\RCRRRCG
		ĺ				RVSEPSPTASWAPAPSLGWAYQ
1		l				LAGGRLVLRVGLSAGWRAACP
l		İ				OGGPISWLEGGLSSGVGLSAG
		ı				WRAACPOGGPISWLEGGLSSST
		l				MRQPRAPQSSLPAPAPWGAPDP
		ı				LAMLAAWIEARPGVWPTOPGP
		ı				wovswvcv
22187	52555	A	22315	11	431	RHLLAAPPPAPPAAAAAAAALSH
1	i	1				CLRSSGRLAPHTSRRLPRVVKR
1	ł					RVNALKNLQVKCAQIEAKFYE
	1					EVHDLERKYAVLYOPLFDKRF
	1	ł				EIINAIYEPTEEECEWKPDEEDEI
						SEELKEKAKIVDEKK\DEEKEDP
		l				KGIPEFW
22188	52556	A	22316	64	467	PAAWLPILVAAROLTVOMMON
					1	POILAALQERLDGLVETPTGYIE
		l				SLPRVVKRRVNALKNLOVKCA
		l		Į.		QIEAKFY\EEVHDLQRKYAVLY
	1	l				OPLFDKRFEIINAIYEPTEEECE
						WKPDEEDEISEELKEKAKIEDE
22189	52557	A	22317	2	177	SSESSISSESICIO INTESE
22190	52558	B	22318	228	377	
22190	52559	A	22319	630	767	WRVKFKNTVKIOANF*KSRFIH
22191	32339	1	22319	0.50	l'*'	LTVVPVTILSOVTLOLTMSPKT
	L	L			L	LIVVEVIILOQVILQLIMSPKI

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide	hnd	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22192	52560	A	22320	80	1362	VTACAAPAAWLPILVADIWSSY
22192	32360	A	22320	80	1302	
		1				NMADIDNKEQSELDQDLDDVE
		1		l		EVEEETGEETKLKARSA*LFR
						MM\QNPQILA\ALQERLDG/LW
		l				VETPT\GYIESLP\RVVKRR\VNA
		1		l		LKNLQVKCAQKETQ\FYE\EVH
		1				DLERKYAVLYQPLFDKRFEIIN
		l				AIYEPTEEECEWKPDEEDEISEE
1	İ					LK\EKGQIEDEKKDEEKEDPKGI
		ı				P*IW\LTVFK\NVDLLSDMVQEH
						DEPILKHLKDIKVKFSDAGQPM
		l l				SFVLEFHFEPNE\YFTN\EVLTKT
		l				YRMRLEPEPDDSDHFSFDGQEI
		l				MGCTGCQIDWKKGKNVTLKTI
		l		i		RK*/RRNHKGRGT\VRTGH*NQ
		l				FPNDSFSNFFCPLLKSPESGDLD
		l				ED\AEAILAADFEIG\HFL\RERII
1		l				PRSVLYFTGEAIEDDDDDDYDEE
		l				GEEADEEGEEGDEENDPDY\D
		l				PKK\DQNPAECKQQ
22193	52561	Α	22321	3	425	GCGCGGAAARGSPAPGRPAAS
		1		ŀ		PPDCPLWLQGCYPPPSCDALSA
		1				TEADGCD*IYPPDTSRDGPHQA
İ		1		İ		SLPGESSSFPGQPNDSYLPERGS
	ł	1		l		ERSRQASYATPAAETQDPDEGL
						POPGPEALPGGLQVPKSAAPFC
						GAQHAAGQ
22194	52562	Α	22322	2	285	SRFEPRVRPTGGDRPHOASPPG
		1				DSSSFPGQPNDSRLPECGSECRG
		l				QASYATPAAETQDPE*RSSPTRS
		l		l		*SP\PGGFOVPKSAAPFCGAOHA
		l		l		AGO
22195	52563	A	22323	85	473	FTMPAVALVPKELYLSSSLKDL
	1	1		I	1	NKKTEVKPEKISTK/ODYVHSIL
1		ı	1	l	I	GTANIKKAIEEAERLSESLKLRY
		l		l		KEAEVRK*LEEKDROKEAOOL
		1		l		OOKROETGRE\DGGTLAKVSLE
		l		l		NVLDYODKTOKSNGEKSEKM
L _						IN A POLICION LOW PROPERSERIAL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop endon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nueleotide insertion)
				sequence		
22196	52564	Α	22324	1	1449	MMRAHMFVYKELKOIYKKKT
			ŀ			HPHOKVGKGYKOTLLRGRHLR
		i				GQETYEKKLTHVYETTDFKQQ
						OYYFHSILGPANIKKATGETER
ŀ						LSESLKLRYEEVEIWKKLEEKD
						RQGEAQWLQQKRQETGREDGS
						MLAKGSLEIVLDSKDKTOKSN
						GEKNEKCETKEKGAITAKELYT
				1		MMMDKNISLIIMDAQRMQDYQ
						DSCILHSLSVPEKAISPGVTASW
						IEAHLPDDSIDTWKKRGNVEY
				l		MVLLDWFSSAKDLQIGTTLWH
			ŀ			LKDALFKWEKGGYKNWFLCY
						SQYTTNAKVTPPPQHQNEELSIS
				i		LDFTYPSLEESIPSKPAAEMPPP
1						PIKVDEDIELISDQISDNDQNER
						TGPLNISIPVESVAASKSDVSPII
						QPVPSIKNVPQIDHTKKLAVKL
1						PEEHIIKSESTNHEQQSPQNEKV
			l			IPDCSTKPVVSSPTLMLTDEEKA
						HIHAETALLMEKNKQEKELQE
						RQQGKQ/K/EKLRREEHEQKAK
22197	52565	Α	22325	185	2983	FIMPAVASVPKELYLSSSLKDL
1			İ			NKKTEVKPEKISTKSYVHSALK
					\	IFKTAEECRLDRDEERAYVLYM
						KYVTVYNLIKKRPDFKQQQAD
						CAYFLGQTVHILFQHRALLEKK
						YRAVGAITAKELYTMMTDKNI
i						SLIIMDARRMQDYQDSCILHSL
1						SVPEEAISPGVTASWIEAHLPDD
1						SKDTWKKRGNVEYVVLLDWF
1						SSAKDLQIGTTLRSLKDALFKW
						ESKTVLRNEPLVLEGGYENWL
						LCYPQYTTNAKVTPPPR
22198	52566	Α	22326	313	3720	FIMPAVASVPKELYLSSSLKDL
	'	1				NKK\TEVKPEKISTKSYVHS/AP
						RKIFKTAEECRLDRDEERAYVL
						YMKYVTVYNLIKKRPDFKQQQ
						DYFHSILGPGNIKKAVEEAERLS
						ESLKLRYEEAEVRKKLEEKDRQ
						EEAQRLQQKRQETGREDGGTL
						AKGSLENVLDSKDKTQKSNGE
				l		KNEKCETKEKGAITAKELYTM
1				l		MTDKNISLIIMDARRMQDYQDS
						CILHSLSVPEEAISPGVTASWIE
1				l		AHLPDDSKDTWKKR
$\overline{}$				L		

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codnn far last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22199	52567	A	22327	46	562	LLOGPGFCGDSMPVEARPSAA
						WRGGRAPGWTPFSPPDRETWK
ĺ						PQ/RASQGESGQRRFGPCPRSGG
ĺ						RPPGPGSAAHSSRSQRWRKASR
						A WPAAPRKGLLPPAQA WPPTP
1						AAGL*GFLAACKR/PPVGGRGP
1						APAPCRPASSRCFLICDSGPPCL
						CSASRAAWMGSAASSRVLFIEG
22200	52568	В	22328	51	445	
22201	52569	A	22329	51	366	CQWNTQTQVCYKQDVQSITAL
				l**		GSAAQCTCLSSPR*SGKFKLLFS
1						PGLPGTLVPSTIPATSAPRA/SHR
1	ļ	l				AGEAWVGEPGRALAALQNSS\P
	ŀ	1				DWAGYOGLPCKPSIRAA
22202	52570	A	22330	257	916	SPSGSNLETHLRAPDLGGPCPP
	ł					GPRWWA*TPQPGHPQTRPSPAA
	1					QPCPFPPQCQQLIHCPC/SPAPGT
						VAAPPAP\PTPASQCLLCPWTSA
						PVCSPAPTFPCPMAPVMAFMLP
						SYSFP/PGTPNLPQAFFPSQPQFP
		l				SHPTLTSEMASASQP\GSPAGPR
		l				SPDSHVLVQPPGPPHHRPWVGP
		l				PHRSFSPAAARPCSSTCCSWRK
						PLRVALEPWGPQGPQRQQL
22203	52571	Α	22331	151	1232	GRLGDARGPQGPEAPGGEEAA
		l				RADSTRACRSCGLLLGGRRGCR
						PSLKNAELLELTVRRVQCVLRG
1		1				RAREREQLQAEARRALRCRLHP
						VACTKVHTFVSTCQAIDATVA
						AELLNHLLESMPLREGSSFQDL LGDALAGPPRAPGRSGWPAGG
						APGSPIPSPPGPGDDLCSDLEEA
						PEAELSOA/PC*GARLGARSPGO
1						PDHSPNCPGVSGGLGDQCQPES
1						CRGGARPSLGSPPLLPGVOMW
1						WG*GPGSLPGSSLPPLMDGLOG
		ĺ				QPLVTSPAQRPSPVS*KNF*GPC
0.		ĺ		ĺ		SSGSGWREGATGRRKNFVELP
						ALSQVSPSPAFNPRLCGLWGQR
						WQKLVLGTSVPGSPGLNKSLN
						LPLORGDEROVH
22204	52572	Α	22332	1	1524	
		-				

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
22205	52573	Α	22333	3	8875	EDGSLLQEVGIICLARGPGCGPE
		İ				TPLYSPATGLALQLQGPAAMG
						ALWSWWILWAGATLLW\GKSG
		l				EDWISGPTTVFSQPGLTQEASV
						DLKNTGREEFLTAFLONYQLA
						YSKAYPRLLISSLSESPASVSILS
		l				QADNTSKKVTVRPGESVMVNI
		l				SAKAEMIGSKIFQHAVVIHSDY
						AISVQALNAKPDTAELTLLRPIQ
		l				ALGTEYFVLTPPGTSARNVKEF
		l				AVVAGAAGASVSVTLKGSVTF
		İ				NGKFYPAGDVLRVT
22206	52574	Α	22334	2	2214	ISATDGPLAPCHGLVPPAQYFQ
ŀ						GCLLDACQVQGHPGGLCPAVA
						TYVAACQAAGAQLGEWRRPDF
						CPLQCPAHSHYELCGDSCPVSC
i	İ				l	PSLSAPEGCESACREGCVCDAG
					ł	FVLSGDTCVPVGQCGCLHDGR
						YYPLGEVFYPGPECERRCECGP
		1				GGHVTCQEGAACGPHEECRLE
						DGVQACHATGCGRCLANGGIH
						YITLDGRVYDLHGSCSYVLAQ
						VCHPKPGDEDFSIVLEKNAAGD
					1	LQRLLVTVAGQVVSLAQGQQV
	l					TVDGEAVALPVAVGRVRVTAE
		ĺ				GRNMVLQTTKGLRLLFDGDAH
						LLMSIPSPFRGRLCGLCGNFNG
					Į.	NWSDDFVLPNGSAASSVETFG
	ł					AAWRAPGSSKGCGEGCGPQGC
		ŀ				PVCLAEETAPYESNEACGQLRN
		ĺ				PQGPFATCQAVLSPSEYFRQCV
					l	YDLCAQKGDKAFLCRSLAAYT
	l					AACQAAGVAVKPWRTDSFCPL
					l	HCPAHSHYSICTRTCQGSCAAL
	İ					SGLTGCTTRCFEGCECDDRFLL
						SQGVCIPVQDCGCTHNGRYLPV
	1					NSSLLTSDCSERCSCSSSSGLTC
	1					Q\AAGCP\PG\RVCEVKAEAR\N
		1	1		l	CW\ATRGL\CVLSVG\ANL\TTF
	1		1			DGARGATTSPGVYELSSRCPGL
			1			QNTIPWYRVVAEVQICHGKTE
						AVGQVHIFFQDGMVTLTPNKG
	1		1			VWVNGLRVDLPAEKLASVSVS
				l		RTPDGSLLVRQKAGVQVWLGA

SEQ ID	SEO IN NO.	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last aminn acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		· · ·
		<u>L</u>	<u></u>			
22207	52575	Α	22335	3	7639	ITCPENSHYEVCGPPCPASCPSP
		1				APLTTPAVCEGPCVEGCQCDA
1		l				GFVLSADRCVPLNNGCGCWAN
1		1				GTYHEAGSEFWADGTCSQWCR
		l				CGPGGGSLVCTPASCGLGEVCG
1		1				LLPSGQHGCQPVSTAECQAWG
1						DPHYVTLDGHRFDFQGTCEYL
		l				LSAPCHGPPLGAENFTVTVANE
		l				HRGSQAVSYTRSVTLQIYNHSL
1		l				TLSARWPRKLQVDGVFVTLPF
		l		l		QLDSLLHAHLSGADVVVTTTS
		1				GLSLAFDGDSFVRLRVPAA
22208	52576	Α	22336	2	224	HCYMVVQLWKPRSLRRKSDQL
		ı				WSSWNQLNTSELSSFISAEMQT
		1				EAFDCGSMAGRM*GQGRYLLA
1		l				ALWDSSVPLS*POAEPWSGVGP
		1				AHLTE/LLSPSRFVSISDLLAPKD
		l				LGTESQVSMEV*KDMGRFHWE
						DIPP*A*VFVIFVFQIFISRTY\DA
		l				TTHFETTCDDIKNLL*R*QGRYL
						LAALWDSSVPLS
22209	52577	Α	22337	2	206	LSSSYLDOPCYETINRIKLYSES
						LARYGKSPYLYPLYGLGELPOG
ļ			1			FAR*EPVFKTSVIQNLIFNIMCI
22210	52578	Α	22338	2	457	
22211	52579	Α	22339	2	1556	FVFPSEVAVGLALSPASIFLSFSS
		l				PFLRAQERLPSLLWWARSPCAF
				ĺ		SLLGSCVRACPAMNEEYDVIV
		l		Ì		LGTG\LTECILSGIMSVNGKKVL
		l				HM\DRNPYYGGESASITALEDL
İ						YKRFKIPGSPPRVKWGEERDW
			1			NVDL\IPKFLKANG\QLVKMLL
		l				YTEVTRYLDFKVTEGSFVYKG
		l				GKIYKVPSTEAEALASSLMGLF
			1			EK\RRFKEFLVYVANFD\EKDPR
į		l				TF\EGIDPKKTTMRDVYKKFDL
	ĺ	1		l		GODVIDFTGHALA\LYRTDDYL\
	l					DOPCYETINRIKLYSESLARYG
		l				KSPYLYPLYGLGELPOGFARLR
		1				CYLWEGTYMLNKPIE\EIIVOE
		1		1		WGKVIGCKI*REEIARCMQLIC
1	l	1		1		DPSYVKDRVEKVGQVITVICIP
1	l	1		l		QPPPSKNTNDANSC\QIIIPQNQ\
	l	ı		1		VNRKSDIYVC\MISFA\HNVAAQ
	l	l				GKSIAIVSTTVETKEPEKEIRPA
1		ı		I		L\ELL\EPIGTEILVSISDLL\VPKD
1		ı		l		LG\TESQIFIS\RTYDATTHFETTS
		l				D\DIKNIYKR\MTGSEFDFEEMK
22212	52580	A	22340	66	420	S. S. H. H. K. K. HOSEI DI BENK
22213	52581	A	22341	64	140	
22214	52582	A	22342	2	254	
	22302	10		1-	·	1

22215 52583 A 22343 82 3530 HILIVAMPEPTKKEENEVPAPAI PPEPSKEKEAGTTPAKDWTL ETPPGEQAKQNANSQLSILE KPQGGTVKVGEDITFIAKVKA DLSEKPTIKWPK/GKWMDLA KAGKHLQLKETFERHSRVYTF MQIIKAKDNFAGNYRCEVTYK DKFDSCSFDLEVHESTGTTPNI IRSAFKRSGEQGEDAGELDFSC LLKRREVKQGEPQVDWEI LKNTKPSEYEKIAFQY/GITDLA GMLKRLKRS	SEQ 1D	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
22215 52583 A 22343 82 3530 HILIVAMPEPTKKEENEVPAPAI PPEPSKEKEAGTTPAKDWTL ETPPGEQAKQNANSQLSILE KPQGGTVKVGEDITFIAKVKA DLSEKPTVKWFKGKWMDLA KAGKHLQLKETFERHSRVYTF MQIIKAKDNFAGNYRCEVTYK DKFDSCSFDLEVHESTGTTPNI IRSAFKRSGEQGEDAGELDFSC LLKRREVKQGEPQVDWEI LKNTKPSEYEKIAFQYGTDLA GMLKRLKRS] GMLRLKRS] GMLRALKRS GAGGOSLHNSC GMLKRLKRS GAGGOSLHNSC GMLKRLKRS GAGGOSLHNSC GMLKRLKRS GAGGOSLHNSC GMLKRLKRS GAGGOSLHNSC GMLKRLKRS GAGGOSLHNSC GMLKALKRS GAGHLQLKETFERHSRVYTE MQIIKAKDNFAGNYRCEVTK GMLKALKRS GAGHLQLKETFERHSRVYTE MQIIKAKDNFAGNYRCEVTK GMLKALKRS GAGHLQLKETFERHSRVYTE MQIIKAKDNFAGNYRCEVTK GMLKALKRSKGEQGEDAGELDFSC LKRREVKQGGEDAGELDFSC LKRREVKQGEPQDVDWE LKNAKPSEYEKIAFQYGTIDLR GMLKALKRREK GAGGOSLHNSC GMLKALKRREK GAGGOSLHNSC GAGGO	NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
22215 52583 A 22343 82 3530		sequence		09/540,217		of peptide sequence	deletion. \=possible nucleotide insertion)
PPEPSKEKEAGTTPAKDWTL					sequence		
PPEPPSKEKEAGTTPAKDWTL ETPPGEQAKQNANSQLSILFII KPQGTVKVQGDITEIAKVKA DLSEKPTIVKWPIK/GKWMDLA KAGKHLQLKETFERISRYVTF MQIIKAKDNFAGNYRCEVTYK DKFDSCSFDLEVHESTGTTPNI IRSAFKRSGEQEQEDGELDPSC LLKREVKQQEEPQVDWEI LKNTKPSEYSKIARQY/GTDLX GMLKRLKRSI TWYRGNPKPKITWMKNKVAT DDPRYMMSNQGVCTLEIRKP PYDGGTYCCKAVIDLGTVEIE CKLEVKVIVQGVNTPGQPVFI EQQQSLINKDF EQQQSLINKDF GCQQSLINKDF CKLEVKVIVQGVNTPGQPVFI EQQQSLINKDF FPPCEPSKEKEAGTTPAKDWTL ETPPGEQAKQNANSQLSILFII KPQGGTVKVGEDITTIAKVKA DLRKPTIKWFKGKWMDLASI AGKHLQLKETFERHSRVYTFE MQIIKAKDNFAGNYRCEVTYK DKPDSCSFDLEVHESTGTTPNI IRSAFKRSGEGEGEDGELDFSC LLKREVKQQEEPQVDWWE LKNAKPSEYEKIAFQYGITDLR GMLKRLKRRREEK GMLKRLKRRREEK FPCCVSLFVVIQLNSDVLPF*P HKKKPTLSHWFLKYHLH-NI NTKALKYRLV*LTSQKH*OM TYPFCNYLKTN*VTNTKVK* EGY*TAKTQKLYPDKVVMAT KPLKTRSPKPAKYGSPHIDAT WTGSGLGLEAS*HSA 22210 52588 A 22349 I 2292 LRHAKWFQARANGLKSCVIVI VLRDLNYTPTWGPLRGWEIL LCEKSIGTANRPMGAGEALR VLECLASGIVMPQHARLARAF	22215	52583	A	22343	82	3530	HLIVAMPEPTKKEENEVPAPAP
							PPEEPSKEKEAGTTPAKDWTLV
							ETPPGEEOAKONANSOLSILFIE
DLSEKPTIKWPIK/GRWMDLA KAGKHLQLKETFERHSRVYTF MQIIKAKDNAGNYRCEVTYK DKPDSCSFDLEVHESTGTIPNI IRSAFRRSGEGEDGAGELDPSC							KPOGGTVKVGEDITFIAKVKAE
KAGKHLQLKETFERHSRYYTE MQIIKAKDNFAGNYRCEVTYK DKFDSCSFDLEVHESTGTTPNI IRSAFKRSGEGGEDAGELDFSC LLKRREVKQGEEPQVDVWEI LKNTKPSEYEKIAFQYGTDLX GMLKRLKRSI			l				
MQIIKAKDNPAGNYRCEVTYK							
DKFDSCSPDLEWHESTGTTPNI IRSAFKRSGGGQEDAGELDFSC LLKRREVKQGEEPQVDVWEI LKNTKPSEYEKIAFQY;GITDLX GMLKRLKRS1 GMLKRLKRS1 DDPRYMENSOQVCTLEIRKP PYDGGTYCCKAVNDLGTVEIE CKLEVKVIYQGVNTPQPVFI EQQQSLINNLOP HLIVAMPEPTKKEENEVPAPAI PPEEPSKEEKAGTTPAKDWTI. ETPPGEQAKQNANSQLSILFII KPQGGTYKVGEDITFIAKVKA DLIKPPTIKWFRGKWMDLASI AGHI-QLKETFERHSRYVTFE MQIIKAKDNFAGKYNTEVTY MGRANDRAGHCPTYK GMLKRLKRIKREEK FOFCVSLFVYIQAISDVLPE*P HKKKPPTLSHWFLK*NLIL*N NTKALKYRKLV*LTSQKH*CQN TYPECNYLKTN-VTTNTKVK* EGY*TAKTQKLYPDKVVMAT KPLKTRSPKPAKYGSPHDTAT WTGSGLGLEAS*HSA 22220 52588 A 22349 1 2299							
IRSAFRRSGEGGEDAGELDFSC LLKRREVKQEEEPQVDVWEI LKNTKPSEYEKIAPQVIGITUL LKNTKPSEYEKIAPQVIGITUL LKNTKPSEYEKIAPQVIGITUL LKNTKPSEYEKIAPQVIGITUL LKNTKPSEYEKIAPQVIGITUL LKNTKPSEYEKIAPQVIGITUL LKNTKPSEYEKIAPQVIGITUL LKNTKPSEYEKIAPQVIGITUL LTVRGNPEKRITVMKINKVAP DDPRYRMFSNQGVCTLEIREP PVDGGTYCCKAVDLOTTVEIE CKLUEVKVIVQCVNTPQOPVT EGQQQSLINKDP LLWAMPEPTKKEEMEVPAPAI PPEPSKEKEAGTTPAKDWTL ETPTGEEQAKQNANSQLSILFII KPQGGTVKVGGEDTITIAKVKA DLLRKPTIKWFKGKWMDLASI AGKHLQLKETFERHSRVYTE MQIIKAKDNFAGNYRCEVTYK DKPDSCSFDLEVHESTGTTPNI IRSAFRRSGEGGEDAGELDFSC LLKRREVKQGEEPQVDWWE LKNAKPSEYEKIAPQVGTDL RGMLKRLKRMRREEK GMLKRLKRMRREEK GMLKRLKRMRREEK GMLKRLKRMRREEK GMLKRLKRMTKEFK-NHLH.**N NTKALKYRLV*LTSQKH*QW TYPFCNYLKTN*VTNTKVK* EGY*TAKTQKLYPDKVVMAT KPLKTRSPKPAKYGSPHDTAT WTGSGLGLEAS*HSA 113 22220 22220 252588 C 22348 47 337 22221 52589 A 22349 1 2292 22222 52589 A 22349 1 2292 22222 52590 A 22350 2 441 LRHAKWFQARANGLKSCVIVI VLRDLUTRYPTWGPLGWELL CKSIGTANRPMGAGEALR VLECLASGIVMPQHAGAGAGATR VLECLASGIVMPQHAGAGAGATR VLECLASGIVMPQHAGAGAGATR VLECLASGIVMPQHAGAGAGATR VLECLASGIVMPQHAGAGAGATR VLECLASGIVMPQHAGAGAGATR VLECLASGIVMPQHAGAGAGATR VLECLASGIVMPQHAGAGAGATR VL		ŀ		1			
LLKRREVKQQEEPQQDVWE LKNTKPSEYEK1AFQY/GITDLX							
LKNTKPSEYEKIAFQY/GITDL\							
Commission			i				
22216 52584 A 22344 66 358							
DDPRYRMENOGOVITLEIREP.	22216	52584	A	22344	66	358	
Pydogtycckaynolgtyee	122.0	52501	ľ.		1		
CKLLEVK VIYQQVNTPQQPVF EGQQQSLHNKDF	i						
EGQQQSLHNKDF							
22217 52585 A 22345 67 3515 HLIVAMPEPITKEENEVPAPAI PPEPSKEKEAGITPAKDWTL ETPPGEQAKQNANSQLSILE KPQGGTVKVGEDITFIAKVKA DLIKRPTIKWPKGWMDLAS AGHI-QLKETFERHSRVYTE MQIIKAKDNFAGWYNGEVTYK MGIKAKDNFAGWYNGEVTYK DKFDSCSFDLEVHESTGTTPNI IRSAFKRSGEQGEDAGELDFSC LLKRREVKQGEPOVDVWE LKNAKPSEYEKIAFQYGITDLR GMIKRILKRIRREEK GMIKRILKRIRREEK FQFCVSLFVYIQLASDVLPEPP HKKKPLYTLSTVKHVW TYPECNYLKTN-VTTNTKVK-EGY-TAKTQKLYPDKVVMAT KPLKTRSPKPAKYGSPHDTAT WTGSGLGLEAS*HSA 22219 52587 A 22347 1 113 22220 52588 C 22348 47 337 22221 32589 A 22349 1 2299 22221 32589 A 22349 1 2299 22222 32590 A 22350 2 441 LRHAKWFQARANGLKSCVIVI VLRDIJVTRVPTWGPLGWPL LCEKSIGTANRPMGAGEALR VLECLASGIVMPQHALRLAF GQLHKVLGMPQHARLAFAF GQLHKVLGMPQHARLAFAF GQLHKVLGMSTENDENSMYKEYP STORT							
PPEPPSKEKAGTTPAKDWTL ETPPGEQAKQNANSQLSILFII KPQGGTVKVQEDITFIAKVKA DLLRKPTIKWFKGKWMDLASI AGKH_QLKEFFERHSRVYTFE MQIIKAKDNFAGNYRCEVTYK DKFDSCSFDLEVHESTGTTPNI IRSAFKRSGEGGEDAGELDFSC LLKRREVKQQEEEPQVDWME LKNAKPSFYEKIAFQVGTDLR GMLKRLKRMRREEK PPEPPSKEKFAGNITAKVAG	22217	52585	A	22345	67	3515	HLIVAMPEPTKKEENEVPAPAP
ETPPGEEQAKQMANSQLSILFII KPQGGTVKVGEDITFIAKVKA DLLRKPTIKWFKGKWMDLASI AGKHI,QLKETFERHSRVYTFE MQIIKAKDNFAGRYKGEVTYK DKRDSCSPDLEVHESTGTTPNI IRSAFKRSGEQQEDAGELDFSC LLKRREVKQQEEPQVDVWE LKNAKPSEYEKIAFQYGITDLR GMIKRI,KRMREEK GMIKRI,KRMREEK FPGFCVSI-FVYIQIASDVLPE*PH HKKKPPTLSHWFLX*NLH.*N. NTKALKYRKI.V*LTSQKH*QN TYPECNYLKTN*VTTNTKVK* EGY**TAKTQKLYVDKVVMAT KPLKTRSPKPAKYGSPHDTAT WTGSGLGLEAS*HSA TEXT							PPEEPSKEKEAGTTPAKDWTLV
KPQGGTVKVGEDITFIAKVKA DLRRPTIKWFKGKWMDLASI		ŀ					
DLIRRPTIKWFRGKWMDLASI		ŀ					
AGKHLQLKETFERHSRVYTFE							DLLRKPTIKWFKGKWMDLASK
MQIIKAKDNFAGNYRCEVTYK							
DKPDSCSPDLEVHESTGTTPNI IRSAFKRSGGQEDAGELDFSC				İ			MOIIKAKDNFAGNYRCEVTYK
IRSAFKRSGEGQEDAGELDFSC		İ		l		1	DKFDSCSFDLEVHESTGTTPNID
LLKRREVKQQEEPQVDVWE		ŀ					IRSAFKRSGEGOEDAGELDFSG
LKMAKPSEYEKIAFOYGITDLR 22218 \$2586 A 22346 2 39 FOFCVSLFYYIQLMSDVLPE*PF							
22218 52586 A 22346 2 39							LKNAKPSEYEKIAFOYGITDLR
HKKKPPTLSHWFLK*NLHL*N: NYKALKYPKLV*LTSQKH*QN							GMLKRLKRMRREEK
NTKALKYRKLV*LTSQKH*OM	22218	52586	Α	22346	2	39	FOFCVSLFVYIOLNSDVLPE*PR
TYPFCNYLKTN*VTTNTKVK* EGY*TAKTQKLYPDKVVMATI KPLKTRSPKPAKYGSPHDTAT							HKKKPPTLSHWFLK*NLHL*N*
EGY*TAKTQKLYPDKVVMAT;							NTKALKYRKLV*LTSQKH*QM
KPLKTRSPKPAKYGSPHDTAT							TYPFCNYLKTN*VTTNTKVK*T
WTGSGLGLEAS*HSA							EGY*TAKTQKLYPDKVVMATN
12220 52587 A 22347 I II 3							KPLKTRSPK\PAKYGSPHDTAT
22220 52588							WTGSGLGLEAS*HSA
22221 52589 A 22349 I 2292		52587					
22222 52590 A 22350 2 441 LRHAKWFQARANGLKSCVIVI VURDLVYRVPTWGPLRGWPL LLCEKSIGTANRPMGAGEALRI VUECLASGIVMPIQHALRLAAF GQLHKVLGMDPLPSKMPKKPR		52588	С	22348	47		
VLRDLAVTR VPT WGPLRG WPL LLCEKSIGTANR PMGAGEALRI VLECLASGIVM PQHALRLAAF GQLHKYLG MDPLPSKMPKK PK	22221	52589	Α	22349	I	2292	
LLCEKSIGTANRPMGAGEALRI VLECLASGIVMP(MALRLAAF GQLHKYLGMDPLPSKMPKKP)	22222	52590	Α	22350	2	441	LRHAKWFQARANGLKSCVIVIR
VLECLASGIVMP/QHALRLAAF GQLHKVLGMDPLPSKMPKKPF					1		VLRDL\VTRVPTWGPLRGWPLE
GQLHKVLGMDPLPSKMPKKPF		1	1	1			LLCEKSIGTANRPMGAGEALRR
							VLECLASGIVMP/QHALRLAAF
		1		1	l		GQLHKVLGMDPLPSKMPKKPK
		1		l	l		NENPVDYTVQIPPSTTYAITPM
KRPMEEDGEEKSPSKKK							KRPMEEDGEEKSPSKKK

SEQ ID	SEQ ID NO:	Mat	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1.01	sequence]"""	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	ļ ·	1		sequence		, , , , , , , , , , , , , , , , , , ,
		 -				
22223	52591	A	22351	45	402	HALAALEKLFPDTPLALDANK
		1				KKRAP\VPVRGGPKFAAKPHNP
		l				GFGMGGPMHNEVPPPPNLRGR
						GRGGSIRGRGRGRGFGGANHG
		l				GYMNAGAGYGSYGYGGNSAT
						QGYK*LFHKTAYGYS
22224	52592	Α	22352	243	3113	FPDTTEEVKMRPMRIFVNDDRH
		l				VMAKHSSVYPTQEELEAVQNM
		l				VSHTERALKAVSDWIDEQEKG
		l				SSEQAESDNMDVPPEDDSKEG
		1				AGEQKTEHMTRTLRG\VMR\VG
						LVAK\CLLLKGDL\DL\ELVLLC\
					İ	KEKPTT\ALL\DKVADNLAI\QL
		1				AAVTEDKFEILQSVDDAAIVIK\
	l		l			NTKEPPLS\LTIHLTSPVVREE\M
						EKVLAGETLS\VNGPPGNVLDR
				ł		SRIAFAAFRGSLPETORWFO\AR
		l				ANGLK\SCV
22225	52593	A	22353	139	245	THIO DAY
22226	52594	A	22354	875	2019	NPNSOMPKMTRSTKOLTNKSG
		1			[· · · ·	OIDGCPGCRVVPTWCRLGHLSH
						YWAVWASILRFWEVWKYWLG
						NMGRKSSEFPILKLNG\RTMGD
		1				*VLPFHMYV\OPVV\HPOLYPHS
		1				VTELTGDYFKPWWDGOPSLOO
		ļ			ļ	VL\ERVDEWM\AKEGLL\DPNV
		1				OVOFLFTLWEDWGLKKSVLPG\
		1				OCO*LGAWPVGGFTSKOWD*S
		ł				*KKAYSF\AMGCWPKNGLLDM
		1				NKGLSLQHIGRPHSGIDDCKNI
		1				ANIMKTLAYGGFIFKOTSKPVLI
1					0	GRGQDGAGQGSLVGPPQNPPL
						PHORGKRESGHKLCCPKVVPSA
	ł					SLWGSMPLGRGLCCPLQGALA
						TWPFLEARHFCAPPSPTPQSQPS
						ISPSHCGPGLRGTPGTHRLLPW
		_				YNRLLWGHQDPRCAPSHP
22227	52595	Α	22355	507	916	RKKNEKEGTKPPRNMGICEKT
1		l				KPTFDWCT*K*RGEWNQVGKY
1	1	1		1	l	TSGYYTGELPQSSKTGQHSNSG
						NTENTTKILLEKSNPKTHNW*I
			1	l		HQG*NEGKTVKGSQREK*GYP
				l		QREAHQTNSRSLCRNPTRQKRV
						GVNIKHS
22228	52596	В	22356	72	359	-

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22229	52597	A	22357	1236	1987	GNSSPFGLPCKSSSSPPPPSLTG APARRAAAPRPASRPRPLAA GGGVPQLFATRPPPJPGPTRARRP TWR*GPRGSPETPRRREWTR REPECCRAQGSRPARPPSGS AVRSASPASSPAPAASAPLGPRS RWS*RPSSRPRNSPPPPPPGSSR SDREWGRKEASPPLTLAEAPA Y/RQDQAPPPGSPHLGFQSAAG
						PSRTASSAP/RSGIGGRGANTPA PAAIFPPSLRPPRAGPVGDLRAS GVDAAA
22230	52598	A	22358	287	2082	IXTETSEEKARRYKDRISQLNA VFQEQKKMIQAQESITLEDVAV DFTWEEWQLLGAAQKDLYRD VMLENYSNLVAVGYQASKPDA LFKIEQGEQUTTEDGIHSGAC SDIWKYDHVLERLQSESLVNRR KPCHENDAFENIVHCKSQFLL GQNHDIFDLIRGKSLKSNLTLLVN QSKGYEIKNSVEFTCNGDSFLH ANNERLHTAIKFPASQKLISTKS QFISPKHQKTRKLEKHHVCSEC KAFIKKSWLTDHQVMHTGER PHRCSLCEKAFSRKFMLTEHQR THTGEKPYECPECGKAFLKKSR LINHQKHTHTGEKPYICSECGKG FIQKGNLIVHQRIHTGEKPYICSEC TPFVCSECGKSCSQKSGLIKHQ RIHTGEKPECSECGKGAFSTKQ KLIVHQKHTHTGERPYICNECGKG TPFVCSECGKSCSQKSGLIKHQ RIHTGEKPECSECGKGAFSTKQ KLIVHQKHTHTGERPYGCNECGK AFAYMSCLVKHKRIHTREKQEF CSQGGKSSCERQALITHO*CHA GEKLC*RGDYTSAFCPSDIIKH QRPHISQNRNCSPCGTASGQM CSLRR**QWICTGQKPCECSECC CAFROQLCLILCYRRYJGKKPJ
22231	52599	С	22359	90	344	
22232	52600	A	22360	2	543	
22233 22234	52601 52602	A	22361 22362	373 111	490 396	GAPPGIWCLSLWACRALSPLSS AAQHRKRPSALSPATPAGSRPL RRCAPGFAATKTQRVQRRPPRR *GAAAGAGGS*RPLCGVVIHSV LHNTPE

SEQ ID		Mct	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		_		1902	3474	Lunana and an an an an an an an an an an an an an
22235	52603	A	22363	1902	34/4	VPQRSTGSGRARSARRPLRAPD
						PCAAAPRVSPQPRPSECSGGRR
						GGSKTWAKRNAGKELTDRNW
					1	DQEDEAEEVGTFSMASEEVLK
		1				NRAIKKAKR\RNVGFESDTGGA
					l	FKGFKRFGGYPSEEERFSGFG\S
		į			i	GAGGKPLEGL\SNGNNIT\SAPP
		ŀ				FASAKAAADP\KVAFGK*LPSPS
			l			RLLVDKNVSNPKTNGDSQQPSS
						SGLASSKACVGNAYHKQLAAL
		l				DCSVRDWMVKHANTNSLCDL
		1		l		TPIFK\DYEK\YLANIETANTGN\
		l			ŀ	SGRNSESESNKVAAETQSPSLF
		l				GSTKLQQESTFLFHGNKTEDTP
		l				DKKMEVASEKKTDPSSLGATS
		l				ASFNFGKKVDSSVLGSLSSVPL
		l			1	TGFSFSPGNSSLFGKDTTQSKPV
		l				SSPFPTKPLEGOAEGDSGECKG
			1			GDEEENDEPPKVVVTEVKEED
				i		AFYSKKCKLFYKKDNEFKEKG
						GTLHLKPTANOKTOLLVRADT
		1				NLGNILLNVLIPP\NMPCTRTGK
						NNVLIVCVPNPPIDEKNA\TMP\
						VTMLIRVKTSEDADELHKILLE
22236	52604	С	22364	65	283	TWENCY RIGEDADEENKIEEE
	52605	A	22365	3	419	
	52606	A	22366	1	512	CGRPPWKTGQRSQPSCPPGSSG
						SGRGIPPAAWHEGEGPEAVLQ
			İ			GHRAGRGDPACRGLCCLPSQL
						GGPTSPTSAASRACGSRGAATL
			İ			VVKVKWFYHP/CRRPSWARGS
						ATARMRCTSPATRMRTTCRPSF
						TSARSWRASSMSRWPGAASAR
						TGRTSTTGGHLRPHHRAPGDG
22239	52607	Α	22367	l .	2940	MRRRKVAAIHAGPRPRSAAA
22239	32007	^	22307	l'	2940	SSEQRGRGPGAARRAGGAEAA
						ASGVTGGGGGGGAGOHAGGLG
		l				FSAGGRAPSORPLCGSMPARLF
		1				
						SKQEVDYCGLPAPPASSLRSRG
		l				HRQRIYSLNQLLQEALRTGRAR
						PGEGPGWLVTDKGGLGWGSSH
			1		1	SQAQGPDSQAAWGRAPPAHGS
		1			1	SPAFLQVLDVRPQSSRYLPPGT
						RVCAYWSQKSRCLYPGNVVRA
		1	l		l	PPPCFGRLLGVPAASSGVPSTG
						ASGDEDEDLDSVVVEFDDGD
	52608	В	22368	215	356	
22241	52609	A	22369	282	499	YLLHCLPSDPPSFTNSPDGRGRF
1 1					İ	RRRQRRRRSDPPGRGRTKSSTR
						VPGSNRFV*FHQR*GRLKEWM ANQDTN

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22242	52610	A	22370	105	619	ACQASSHSGPVWGPGERTTPG A*ASARREPSRNQHPNLWVTE Q1*IASPRTVANPLVPARTPTGP RTNKPKHLIKQGRQPRQPLSG STGGAGESPASGCAGGQPAHR VWIAPAARGCQHTPWKAPAG MQQSPRPASATRMIGPSPLSP APPAVVGVAALGSRNRCIL
22243	52611	A	22371	1	441	
22244	52612	A	22372	10	621	LRQLERGLAKGQPSTPVDRPLG PDPSAPGTLAGAALPPLEPPAPC LCQDPQEDSVEDEEPPGSLGLP PPQAGVQPAAAAVSGTTQPL AGPOVQPAAAAVSGTTQPL AGPOVQPAAAAVSGTTQPL AGPOVQPAAAAVSGTQP AGPQGLSAPEQQEDEDSLEEDS PRALGSGQPABSAELDE QDILAPQTVQCPAQAPAGGSEE TLRQLERGLAKGQPSTPVDRP LCPDPSAPGTLAGAALPPLEPP APCLCQDPQEDSVEDEEPPGSL GLPPPQAGVQPAAAAVSGTTQP LGTGPRVSLSPHSPLLSPKVAS MDAKDLALQILPPCQVPPPSGP QSPAGPQGLASPEQGEDSLEEDS EDSPRALGSGQHSDSHGESSAE LDEQDILAPQTVQCPAQAPAGG SEETIAKA

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SEQ ID	SEQ ID NO:	Mct	SEQ ID NO:		Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22245	52613	A	22373	I	2271	MPGELLAPSCCCRGGPTRDRTD
22243	52015	ľ^	22313	l'	1271	LSCDAAAATTILGGDRREPCAL
		1				TPGPSHLALTFLPSKPGARPOPE
		1				GASWDAGPG\LSTEAPLPATLE
		l				
						PRIVMGEETCQALLSPRAARTA LRDOEGWGL/SPOGSMVDEREL
		l		İ		HPAGTPEPPSSESSLSADSSSSW
						GOEGHFFDLDFLANDPMIPAAL
		ŀ				
						LPFQGSLIFQVEAVEVTPLSPEE
		1		ŀ		EEEEAVADPDPDSASYAEADDE
		ŀ		l		RLYSGEPHAQATLLQDSVQKTE
						EESGGLRYDTASAPTRHRSLVS
i						PKACSCSHDCAP\AG*RGPHLTP
						GLRYDTASAPTRHRSLISPKAC
					ŀ	GCSHDCVPAG*RGPHLTPGLRY
		1	İ			DTASAPTRHRSLVSPKACSCSH
1		1				ACVPAG*RGPHLTPGLRYDSTS
-		l	1			ASARHRPHLRSRASGCGHPSNL
1		1			1	AGRSRLCPRDRACGHHGSAGS/
		1			ŀ	SAWTGSMSGSA\RHR*WG*A*L
		ı				TPKGDPGG*EPAGRRPEG\TGT
ıl .						GTWTQPQGGLHGCQRPGLADL
	1					AALPSASSLWAPEPSWPS/SALS
		ł				PRAARG*GQPGGRLASGPGLGP
		l				AFG*PRGVISRAGRAGHLGSSD
		1			1	RAVSSPGAQSREEGPSRRQ*GD
		1				HRQSQAESQ*EEGPKGNVKAG
		l				LAADSGSHQDHHPEVQEHPLC
1		l				HRQA*CLQGDPASAHLCG\FWR
1						GQ\T*GP\CPASAQSRS*EV*GAL
		l				R\DLSLGP*VST\QGPG*A*VQG
		1	l			KGKG\GGGKGGPGGAGTA*HL
22246	52614	Α	22374	2	211	
22247	52615	Α	22375	43	241	
22248	52616	A	22376	184	332	
22249	52617	Α	22377	1292	1923	KAGARGNQEGTMASSSG\TQK
		1				VLTIPRGCYQ*K*SKETLP*CSG
		i				WPNDAR\EAVVNC\CTEFI\HLIS
						SEANEICNKSEKKTISPEHVIQA
		l				LESLGFGSYISEVKEVLOECKT
1		ı				VALK\RRKASSRFGKPLGIPEEE\
1	1		l		1	LFGDQANKELFGKSLDQQQAE\
				1		LAQQ\EW\LQMQQAA\HQA\QL
1	1		l		1	AAASAKCI*SRRGSFSGMEEDG
			ĺ	1		GWISEIPPAEFLFLL
22250	52618	В	22378	1	1701	
22251	52619	A	22379	1	147	
22231	1520.5	٠.	1223.3	l'	1	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
22252	52620	A	22380	1	484	IRHEGRGQRGKMELVQ\VLKRG
		l	İ			LQQITGHGG/LFRGYLRVFF\RT
		l				NDAKVGTLVGEDKYGNKYYE
		ı				DNKQFFGRHRWVVYTTEMNG
		ĺ				KNTFWDVDGSMVPPEWHRWL
		ı				HSMTDDPPTTKPLTA\RKFHLG
		l				RNH*ISTWTGAPQEQYVPYSTT
						RK\KIQEWIPPSTPYK
22253	52621	Α	22381	3	258	DMAAQIAEGMAFIEERNYIHRD
		l				LRAANILVSDTLSCKIADFGLA
		l				RLIED\FTIKSDVWSFGILLTEIV
						THGRIPYPGMTNPEVIQN
22254	52622	Α	22382	1	738	
22255	52623	Α	22383	233	1898	GAGPTGYLRRLPGWAGTMGC
		1				GCSSHPEDDWMENIDVCENCH
		1				YPIVPLDGKGTLLIRNGSEVRDP
						LVTYEGSNPPASPLQDNLVI\AL
					!	HSYEPSHDGDLGFEKGEQLRIL
						EQSV\EWWKAQSLTTGQEGFIP
		l				FNFVAKANSLEPEPWFFKNLSR
i						KDAERQLLAPGNTHGSFLIRES
		1				ESTAGSFSLSVRDFDQNQGEVV
		l				KHYKIRNLDNGGFYISPRITFPG
		l			1	LHELASAITPIASDGLCTR\LSRP
						CQTQKPQKPQKPWWEDEWEV
		l				PRETLKL VERLGAGQFG\RCGW
		l				GTTTGTTKVAVKSLKQGSMSA
		1			İ	GRLP\AEANLMKQLQ\HQRL\VR
		l			1	LYAVVT\QEPIYIITEYMENGSL
	ļ	l				VDFLKTPSGIKLTINKLLDMAA
		l			1	QIAEGMAFIEERNYIHRDLRAA
						NILVSDTLSCKIADFGLARLIED
		l				NEYTAREGAKFPIKWTAPEAI*S
	l	l				RGNSPPRSDVWSF\GDLLPEIFT\
1		l				HGRIP/YPQGIPNPELIQKLERGY
		1				RMVRP\SNCPEKLYQL\MRLCW
	l	l			1	KERPED\RPTFDYLRSVLEDFFK
		l				RHRRASNQPQALREPLRKPLCF
22256	52624	Α	22384	3	203	
22257	52625	Α	22385	2	467	
22258	52626	Α	22386	1	468	
22259	52627	A	22387	1	645	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22260	52628	A	22388	821	2637	AAASAWRATHSTTAAGRRTAL
						FRSPEPGNTRDTRRRRRATAPD
ŀ		l				CCRPATSPRALDWTSLHTRRHP
1						VQNYLQWSYWQVLVSRWIAF
l						AGLAMADYLISGGTGYVPEDG
						LTAQQLFASADDLTYNDFLILP
l						GFIDFIADEVDLTSALTRKITLK
						TPLISSPMDTVTEADMAIAMAL
ł						MGGIGFIHHNCTPEFQANEVRK
						VKNFEQGFITDPVVLSPSHTVG\
				1		DVLEGKMRHGFSG\IPITETGTM
				1		GSKLVGIVTSRDIDFLAEKDHT
i i				1		TLLSEVMTPRIELVVAPAGVTL
İ					l l	KEANEILQRSKKGKLPIVNDCD
					ļ	ELVAHARTDLKKSRDYPLASK
						DSQPQLLCGAAVDTREEDKYR
1		l				LDLLTQAGVDVIVLDSSQGNSV
		Į.			i	YQY\ARVLYFKQKYPHLQV\LG
		1		ŀ		GNVVTAAQAKNLIDAGVDGLR
		l				VGMGCGSICITQ\EVMACGRPQ
		l				GTCCVTR*AEYARRFGVPIIAD
		1				GGIQTVGHVVKALALGASTVM
		i				MGSLLAATTEAPGEYFFSDGVR
		l				LKKYRGMGSLDAMEKSSSSQK
		l				RYFSEGDKVKIAQGVSGSIQDK
		l				GSIQKFVPYLIAGIQHGCQDIGA
	ŀ	l				RSLSVLRSMMYSGELKFEKRT
	į .	l				MSPQIEGGVHGLHSYEKRLY
22261	52629	Α	22389	2	332	
22262	52630	Α	22390	3	608	MTAVHAGHLNLKWDPKSLEIR
		1				TLAAERLVTQVTTLVNTNSKGP
		l				SDKKRGHSKKVHVLAASVEQA
		1				AENFLEKEDKIVKESQF\L*E*LL
İ		i				AAVEYVRKQ/GKECLRPSLEEH
	ł	1				LEGIISGAALMGDSSCKHDDHR
		ĺ			1	ERIAAECNTVLQ\ACRTCF/SEY
	1				1	MGNAGCKERSNSLNSAIDKMT
		1			1	KKTRDLRRQLHKAVMDHVSDS
	1	l				FLETNVPLL
22263	52631	A	22391	2	2864	

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \-possible nucleotide insertion)
22264	52632	Α	22392	14	2897	PDWRETKQRPSASGLWNLALA
		1				QLAAEMTAVHAGHLNLKWDP
		1		İ		KSLEIRT\LAVERLLEPLVTQVT
i						TLVNTNSKGPSDKKRGHSKKV
		l				HVLAASVEQAAENFLEKEDKIV
l	l	ı				KESQF\L*E*LLAAVEDVRKQG
		1				DLMKAAARRRSHDDPCSSVEQ
		ı				GNMVQAA*TLLSAVTCLLILA\
		l	ŀ			DMGNVYKLLVQLKVVEDGILK
						LRNAGNEQDLGIQYKALKPEV
		1				DKLNIMAAKRQQELKDVGHRD
		L_				QMAAARGILQKNVPILYTAS
22265	52633	Α	22393	1	1584	
22266	52634	Α	22394	I	3519	
22267	52635	Α	22395	3	673	
22268	52636	Α	22396	I	1574	DVPRDLEVVAATPTSLLISWDA
		l				PAVTVRYYRITYGETGGNSPVQ
		1				EFTVPGSKSTATISGLKPGVDYT
	1	l		'		ITVYAVTGRGDSPASSKPISINY
	l	l				RTEIDKPSQMQVTDVQDNSISV
		l			1	KWLPSSSPVTGYRVTTTPKNGP
l		l				GPTKTKTAGPDQTEMTIEGLQP
		l			ŀ	TVEYVVSVYAQ\NPSGESQPLV
		l				Q\VAVPTIPAPTDLKFTQVTPTS
		l				LSAQWALPNVQLTGYRVRVTP
		l				KERTGPMKEINLAPDSSSVVVS
		l				GLMVATKYEVSVYALKDTLTS
	l	1			l	GPAQGVVTTLENVSPPRRARVT
		l			İ	DATETTITISWRTKTETITGFQV
		l			1	DAVPANGQT\PIQRTIKPDVRSY
	1	l				TITGLQPGTDYKIYLYTLNDNA
		l				RSSPVVIDASTAIDAPSNLRFLA
	1	l				TTPNSLLVSWQPPRARITGYIIK
	1	ı				YEKPGSPPREVVPRPRPGVTEA
	1	I			1	TITGLEPGTEYTIYVIALKNNQK
	1	I			1	SEPLIGRKKTVQKTPFVTHPGY
	1	ı				DTGNGIQLPGTSGQQPSVGQQ
	1	I				MIFEEHGFRRTTPPTTATPIRHR
		L				PRPYPPNVGQEALS

SEQ ID	SEQ ID NO: of peptide	Met	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	sequence	noa	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22269	52637	Α	22397	3	2382	QTEMTIEGLQPTVEYVVSVYAQ
						NPSGESQPLVQTAVTNIDRPKG
						LAFTDVDVDSIKIAWESPQGQV
					İ	SRYRVTYSSPEDGIHELFPAPDG
						EEDTAELQGLRPGSEYTVSVVA
						LHDDMESQPLIGTQSTAIPAPTD
		i			i	LKFTQVTPTSLSAQWTPPNVQL
					l	TGYRVRVTPKEKTGPMKEINLA
						PDSSSVVVSGLMVATKYEVSV
						YALKDTLTSRPAQGVVTTLEN
						VSPPRRARVTDATETTITISWRT
	1				1	KTETITGFQVDAVPANGQTPIQ
					1	RTIKPDVRSYTITGLQPGTDYKI
						YLYTLNDNARSSPVVIDASTAI
		1				DAPSNLRFLATTPNSLLVSWQP
		l				PRARITGYIIKYEKPGSPPREVV PRPRPGVTEATITGLEPGTEYTI
					1	
		1			1	YVIALKNNQKSEPLIGRKKTDE LPQLVTLPHPNLHGPEILDVPST
						VOKTPFVTHPGYDTGNGIOLPG
	1				1	TSGQQPSVGQQMIFEEHGFRRT
ľ	l					TPPTTATPIRHRPRPYPPNVGEE
						QIGHIPREDVDYHLYPHGPGLN
	1					PNASTGQEALSQTTISWAPFQD
ĺ					1	TSEYIISCHPVGTDEEPLOFRVP
	i					GTS\TSATLTGLTRGATYNIIVE
						ALKDQQRHKVREEVVTVGNSV
	1					NEGLNOPTDDSCFDPYTVSHYA
İ	1	1				VGDEWERMSESGFKLLCOCLG
				1		FGSGHFRCDSSRWCHDNGVNY
		1			l	KIGEKWDRQGENGQMMSCTCL
1				1		GNGKGEFKCDPHEATCYDDGK
22270	52638	Α	22398	48	7500	GATMACCAAQGEERVTPCASG
						KRGPAATTSLVLWIPSVPPPVPF
	1					PTLWPPPSWRRQPPGGIRRDFS
				1		RRLRREANLVATCLPVRASLPH
						RLNMLKGPGPGLLLLAVLCLG
1				1	1	TAVPSTGASKSKRQAQQMVQP
				İ		QSPVAVSQSKPGCYDNGKHYQ
}						INQQWERTYLGNVLVCTCYGG
1						SRGFNCESKPEAEETCFDKYTG
						NTYRVGDTYERPKDSMIWDCT
				ŀ		CIGAGRGRISCTIANRCHEGGQS
						YKIGDTWRRPHETGGYM
22271	52639	Α	22399	1	403	NSTNKRRGFCFITFKE\EMEKKY
			l			HNVGLSKCEIKVAMSKEQYQQ
						QQQWGSRGGFAGRARGRGGG
			l		i	PSQNWNQGYSNYWNQGYGNY
	1		İ		ŀ	GYNSQGYGGYGGYDYTGYNN
			l			YYGYGDYSNQQSGYGKVSRRG
	1		ı			GHQNSYKPY

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence	liou	09/540,217	codon for peptide sequence	of peptide sequence	deletion, =possible nucleotide insertion)
22272	52640	С	22400	36	214	
22273	52641	Α	22401	3	182	
22274	52642	Α	22402	223	452	LESMTIFPYSDDKRAQYIFIQKD
						LLKKPVRHYYIMTLSHWAFNT
1		ł				RLALQYWGKGHKTQIDSEA*Q
		ļ				LRNKTMKANFF
22275	52643	Α	22403	1	1037	MSEEQFGGDGAAAAATAAVG
l						GSAGEQEGAMVAATQGAAAA
1		1				AGSGAGTGGGTASGGTEGGSA
		1		1		ESEGAKIDASKNEEDEGHSNSS
						PRHSEAATAQREEWKMFIGGLS
		1				WDTTKKDLKDYFSKFGEVVDC
İ		1				TLKLDPITGRSRGFGFVLFKESE
		1				SVDKVMDQKEHKLNGKVIDPK
		1		i		RAKAMKTKEPVKKIFVGGLSP
		1				DTPEEKIREYFGGFGEVESIELP
		l				MDNKTNKRRGFCFITFKEEEPV
		l				KKIMEKKYHNVGLSKCELK\VA
		1				\MSKEQYQQ\Q\QQWDSRGGCA
		1				GRA\RGRGGDQAE\GYGKGIQA
		l				RWAIKNSYKPYLNYSICQPYPP
						TAGTSLQALCRADFRFSQARSM
22276	52644	Α	22404	299	1508	HYVGGSSSGGDGAAARAT/AR
		l				AVGGSAGEQ\EGA\MVAATHG
i		l	,			ARGGVWEADAATGGGTASGG
		l				T\EGGSAE\SEG\AKIYASKNEED
		l				EGHSNSSPRHSEAATAQRGRM\
		l				KMVIGGLSWDT\TKKDLKDYFS
1		l	ŀ			KFGEVVDCTLKLDPITG\QSRGF
		l				GFV\LFKESESV\DKVMDQKEH
1		l				K\LNG\KVNDPKRAKAM\KTKE
		l		l		PVKKIFVGGLSPDTPE\EKIREYF
		l				GGFGE\VESIELPMD\NKTN*RG
		1		ļ		RGF\CF\ITFKEEEPVKKI\MEKK
		1		l		YHNVGLSKCEIKVVLLNEQYQ
		l				QQQQW*SIGGFAGIALVIGGGP
		l				SQNWNQGYSNYWNQGYGNYG
		l	l			YSS\QGYG\GYGGYDYTGY\NN
	1	1	1			Y\YGYGDYSNQQSGYGKVS\RR
1		l				GGHQ\NSYKPHTKFIPICNLFPN
		L.				RWLKQYFPNLKIQFERGSCHLL
22277	52645	Α	22405	3	189	EESEPERKRARTDEVL\PEEAAP
		1			1	RRKMRTTRTTCPV\$RY\AAP\QL
L		_				LLQKLLQRRRKGSCSRG
22278	52646	В	22406	1	1356	
22279	52647	С	22407	135	260	
22280	52648	A	22408	1	1740	
22281	52649	В	22409	80	371	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22282	52650	A	22410	257	1200	ISWCCMEQKKRLP\FSKREG\PY
	1					EFIICPSEDSSPLLRCAFCIGGMS
						VKEQMETIRHPLQTTRGPGPGP
						LFTDGPDPIPEAFLHGLLSIREM
						VSTEKVKPVGQESEPRVKPLEL
						DDSASGGVHMMVATPGRLMD
ì						LLQKKMVSLNICRYLALDEAD
						RMIDMGFEGDIRTIFSYFKHTSA
						AAIGLQLALDGGTGSGQKGKC
						EEFPSSMSAVSTGLEASACCLS
						VTMDPLETLSRMASSTSSMSSL
					l	ARAWLSPEPGDSPGMVGGLPR
						ELPGMAPLECGEFQRPESLCAD
						RNELGSQSPEFPGQEEEDEDED
						DEEEDEDE
22283	52651	A	22411	107	2069	GGVSPWRACVQQRMEESEPER
						KRARTDEV\LPEEAAPRRKMRT
						TRTTCPMCRYGSAGSLLLQKLL
						QR\RRKGAAEEEQQDSGSEPRG
						DEDDIPLGPQ\SNVSLLDQHQHL
		ŀ				\KEKAESAAKESCQGRKQLEGK
						KEKILGRIVCRRAE\LDVK*KAD
						WLKGITVLIDPIKTQLGLPPR/Y
			i			MFLSMSEERHERVRKKYHILVE
				ł		GDGIPPPIKSFKEMKFPAAILRG
İ				1		LKKKGIHHPTPIQIQGIPTILSCR
			ì			DMIGIAFTGSGKTLVFTLPVIMF
						CLEQEKRLPFSKREGPYGLIICP
						SRELARQTHGILEYYCRLLQED
						SSPLLRC\ALCIGGMSVK\EQ\ME
						TIRQGVHMMVATPG/RALMDL
						LQKKMVSLDICRYLALDEADR
ĺ						MIDMGFEGDIRTIFSYFKGQRQ
						TLLFSATMPKKIQNFA\KSALVK
						PVTINVGRAGAASLDVIQEVEY
						VKEEAKMVYLLECLQKTPPPV
			İ			LIFAEKKADVDAIHEYLLLKGV
	ĺ					EAVAIHGGKDQEERTKA\IEAFR
		1	l	l		EGKKDVLVAT\DVASKGLVFPA
						IQHVINYDMPEEIENYVHRIGRT
1				l		GR\SGNTGIATTFINKACDESFLI
				l		D\LKALLVRKPKQKVPPVL\QV
1			1			L\HCGGWSPMLDIGRSSRGCAF
		l				CGG\LGH\RITDCPKL\EA\MQTK
		L				QVSNIGRKDYLAHSSMDF
22284	52652	A	22412	435	1428	

	possible musteritate ble nucleotide insertion) LVTRLQKALGVRQ RAKMAMSHEPBN SKNNIVCKRLNWP GHLDDPASQEIEQ WPDHVAMLQFISKI NHRMRRYLSK/TG EFQDHLVPDFGCH ELKPHINGPFTPD GKVAEKEGWPLDI
1 22285 52653 A 22413 3 3 597 HKMAPYSPI YHVASVLCC EYIHVDLE LTILLERINY GK/S/YLWL GATISVFPY RTDIANLAD YDQLIEINLS QAHPVAEVC RYCLT	DIE nucleotide insertion) VTRLQKALGVRQ RAKMAMSHFEPN RKNNIVCKRLNWP GHLDDPASQEIEQ WPDHVAMLQFISKI. WHRMRRYLSK/TG EFQDHLVPDPGCH EELKPHINGPFTPD
22285 52653 A 22413 3 597 HKMAPYSPI	ORAKMAMSHFEPN KNINIVCKRLNWP GHLDDPASQEIEQ WPDHVAMLQFISKI NHRMRRYLSK/TG EFQDHLVPDPGCH SELKPHINGPFTPD
YHVASVLCC	ORAKMAMSHFEPN KNINIVCKRLNWP GHLDDPASQEIEQ WPDHVAMLQFISKI NHRMRRYLSK/TG EFQDHLVPDPGCH SELKPHINGPFTPD
YHVASVLCC	ORAKMAMSHFEPN KNINIVCKRLNWP GHLDDPASQEIEQ WPDHVAMLQFISKI NHRMRRYLSK/TG EFQDHLVPDPGCH SELKPHINGPFTPD
EYIHYOLLE LTLLEKIVY0 GKSNYLWL GKSNYLWL GKSNYLWL GKSNYLWL GKTSNYEW RTDIANLAD YDQLIEINLS QAHPWAEV0 RVGLT	KNINIVCKRLNWP GHLDDPASQEIEQ WPDHVAMLQFISKI NHRMRRYLSK/TG ÆFQDHLVPDPGCH EELKPHINGPFTPD
LTLLEKIVY GK/S/YLWL/ GATTISVFPY RTDIANLAD YDQLIEINLS QAHPVAEV RVGLT 22286 52654 A 22414 29 512 22287 52655 A 22415 1 543 22288 52656 B 22416 271 719 22289 52657 A 22417 1 2438 SVIKMAPY: RQYHVGSVI	GHLDDPASQEIEQ WPDHVAMLQFISKI NHRMRRYLSK/TG EFQDHLVPDPGCH SELKPHINGPFTPD
GK/SYLWAL GATISVFPY RTDIANLAD YDQLIEINLS QAHPVAEVG RVGLT 1 543 22288 \$2656 B 22416 271 719 22288 \$2657 A 22417 1 2438 SVIKMAPYS RQYHVGSVI	VPDHVAMLQFISKI NHRMRRYLSK/TG EFQDHLVPDPGCH SELKPHINGPFTPD
GATTSVFPY RTDIANLAD YDQLIEINLS QAHPWAEVC RVGLT	NHRMRRYLSK/TG EFQDHLVPDPGCH EELKPHINGPFTPD
GATTSVFPY RTDIANLAD YDQLIEINLS QAHPWAEVC RVGLT	NHRMRRYLSK/TG EFQDHLVPDPGCH EELKPHINGPFTPD
VPQQLEINLS QAHPVAEVG RVGLT	ELKPHINGPFTPD
VPQQLEINLS QAHPVAEVG RVGLT	ELKPHINGPFTPD
QAHPVAEVC RVGLT 22286 52654 A 22414 29 512	
RVGLT RVGLT RVGL	SKYNDKOO WI EDI
22286 \$2654 A 22414 29 512 22287 \$2655 A 22415 I 543 22288 \$2656 B 22416 271 719 22289 \$2657 A 22417 I 2438 SVHKMAPY-RQYHVGSVI	
22287 52655 A 22415 I 543	
22288 52656 B 22416 271 719	
22289 52657 A 22417 I 2438 SVHKMAPY: RQYHVGSVI	
RQYHVGSVI	SLLVTRL*KALGV
	YGHLDDPASOEIE
	PDRVAMQDA\TA
	SSGLSKVAVPSTIH
	GGEE\DL\RRAKDI
	TAGAKYGVGFW
	ILENYCVPWLSF*L
	GG\LGGI\CIGSCG
	NGLGIPWELKCPK
	LSGWPSPKDVILK
	GTGAIVEYQGPGV
DSISCTGMA	TICNMGAEIGATT
SVFPYNHRN	IKKYLSKTGREDIA
	ILVPDPGCHYDQLI
EINLSELKPI	IINGPFTPDLAHPV
AEVGKVAE	KEGWPLDIR\VGL\I
GSCTNFKLW	/KIWGRSAAVAKQ
	KSQF\TITPGFPRQI\
RATFE\RDG\	YAQILRDLGG\IVL
ANACGPCIG	QWDRKDIKKGEK
	NFTG\RNDANPET
	TALAIAGTLKF\N
	GWARSFRLGGSRI
	F*PRGRDTYOHPP
	WDLSPTSQRLQLL
	KDLEDLOILIKVKG
	AGPWLKFRGHLD
	INIENGKANSVG\
	APD\LARYYKKHG
	NYGEG\SSREHAA
	SPVCGP/CSAGGGP
	ESSSVPRGG/CCSP
	CFPK*SDGSGCAE
APDSGE/CS*	ALGPHCGD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hnd	09/540,217	codon for peptide sequence	of peptide sequence	*"Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22291	52659	A	22419	160	460	RLCIGDAGVSPVCGPLSAGGGP R*LNL*GGEESSSVPRGG/CCSP
						VTATTAPGPCFPK*SDGSGCAE
	ŀ					APDSGNCGAFCALRSPHRHPGP
						KPARERLLLTGL
22292	52660	Α	22420	43	418	SLGCAEQN/FPAIAIQPGMAQEE
	İ					RLSRYQQFKDFQRRILVATNLF
						GRGMDIERVNIVFNYDMPEDS
						DTYLHRVARAGRFGTKGLAITF
					1	VSDENDAKILNDVQDRFEVNV
						AELPEEIDISTYIEQSR
22293	52661	Α	22421	I	592	RFSKYMPSVKVSVFSRGLSIKK
						DEEVLKKNCPHVVVRTRARILA
						LVRNRSFSLQNVKHFVLDECD
						KMLEQLDMRRDVQEIFRLTPHE
						KQCMMF/ERLSRYQQFKDFQRF
ì						ILVATNLFGRGMDIERVNIVFN
						YDMPEDSDTYLHRVARAGRFG
					l	TKGLAITFVSDENDAKILNDVQ
						DRFEVNVAELPEEIDISTYIEQS
22294	52662	Α	22422	2	575	
22295	52663	В	22423	100	412	
22296	52664	Α	22424	2	441	
22297	52665	Α	22425	682	1032	
22298	52666	Α	22426	I	1324	MAEQDVENDLLDYDEEEEPQA
						PQESTPAPPKKDIKGSYVSIHSS
		İ				GFRDFLLKPELLRAIVDCGFEHF
	1					SEVQHECIPQAILGMDVLCQAK
	1					SGMGKTAVFVLATLQQIEPVN
			ł			GQSGQVHIFGLHIGARALVISEH
						ILKDEGAVEGCGVVQCLMARV
	ŀ					TVLVMCHTRELAFQISKEYERF
						SKYMPSVKVSVFFGGLSIKKDE
	l		ļ			EVLKKNCPHVVVGTPGRILALV
	1	1				RNRSFSLKNVKHFVLDECDKM
1		1	l	I		LEQLDMRR\DVQEIFRLDTTPR
				l		KQCMMFSATL\SKDI\RPV\CRK
1				l		FMQDPMEVF\VDDETKLTLHG\
		1	I	I		LQQYYVKLKTSEKNRSLFDLL\
						DV\LEFNQV\IIFVKSVQRCMAL
		1	I	i		AQLL\VEQNFPAI\AIHRGMAQE
			İ	l	1	ERLSRYQQFKDFQRRILVATNL
	1	1	l	I		FGRGMDIERVNIVFNYDMPEDS
	1	l	I	I	1	DTYLHRVARAGRFGTKGLAITF

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22299	52667	A	22427	9	1535	EAQQLVSERPAENRSWKCLLA ARGEERGASIMAEQDVENDLL DYDEEEEPQAPQESTPAPPKKD IKGSYWSIHSSGFRDF/LCLKPE LLRAIGGLVAFEHPSEVQIECCI PQGHPGAWDVLLPGQSSGMCK TAVFRCWPPLQNEPVNGQVT LVMCHTRELAFQISKEYERS KYMPSVKVSVFFGGLSIKKDVE EVMKKNCPHVVJGPTGRILAL VRNRSFSLKNVKHFVLDECDK MLEQLDMRRDVQEIFRLTPHEK QDPMEVFVDDETKLTLHGLQQ YYVKLKDSEKTRKVFDLWDV LEFNQVDLRQGQCKRCMALA QLLIVEQNIFPAGWPHIRGNGFQ VATQILIFGIRGMDIEAKSINIVEN VAMPFELGETVLHRVARSGR EGTKRPPAFTFVSDGNDAIKILN
	50.00		22.422		200	DV\QDRF*S*MWAEL\PEEIDIST YIEQSRLNHQLARAGPPGAAPH AASPPLSRRHC
22300 22301	52668 52669	A	22428	3	290 690	MNQTSEKEIISKEQGNVKAAGA SDVVLYKIDVPANRYDLLCLEG LVRGLQVFKERNFFLTVLEVEK SKTKVEQIQCLCSQEDIADKLG VDISATKAVHISNPKTAEFQV KTILLFGLEKTIAANRKMPLA KITLLFGLEKTIAANRKMPL KLFEISDIVIKDSNTDVGAKNVR HLCAVYYNKNPGFEIHGLLDRI MQLLDVPPGEDKGGYVIQSVG KLGVLHPDVITKFELTMPCSSL EINIGPFL

2900

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22302	52670	A	22430	2	1834	GTRPTVSVKRDLLFQALGRTYT
22302	52070	ļ.,	22.50	ľ	1.05	DEEFDE\LCFEFGLELDEITSEKE
		i				IISKEOGNVKAAGASDVVLYKI
1		1		l		DVLATRYDLLCLEGLVRGLOV
		l		ĺ		FKERIKAPVYKRVMPDGKIOKL
		l		ŀ		IITEETAKIRPFAGAAVLRNIKFT
		1		İ		KD\RYDSFIELQEKLHQNICRKR
		l		ŀ		ALVAIGTHDLDTLSGPFTYTAK
		1		ł		RPSDIKFKPLNKTKEYTACELM
		1		ł		NIYKTONHLKHYLHIIENKPLYP
		1		l		VIYDSNG\VVLSMPPIINGDHSRI
		1		l		TVNTRNIFIECTGTDFTKAKIVL
l		l				DIIVTMFSEYCENQFTVEAAEV
		l				VFPNGKSHTFPELAYRKEMVR
ŀ				l		ADLINKKVGIRETPENLAKLLT
ŀ				i		RMYLKSEVIGDGNOIEIEIPPTR
				l		ADIIHACDIV\EDAAIAYG\YNNI
		l		l		\QMHSPENFTP*ANQFPLNK\LT
		ı				ELLRHDMA\AAGFH*STLPLAL
		l		İ		CSQGRYWLDKLGVGYLWQQR
		l				AVHISN\PKTAEFQVARTTLLPG
		l		l		PP*RP*AANRKV\PLPL\KLFEIS\
		l		i		DIVIK\DSNT\DVGAKKTTROSL
		ı				VAGLLPTKNPG\FEINSWGCWD
	ĺ	l				RIMAVCSDV\PPGGRQGGDM*F
		1				KASEGPAFFPGAMCRDSFARGQ
		ł				SL\GKLG\VLHPDVIHQIWS*PLP
		ł				CSSLEI\NIGPFL
22303	52671	A	22431	I	304	CSSLEIMIGFFE
22304	52672	A	22432	1	642	
22305	52673	В	22433	1	423	
22306	52674	A	22434	i	525	LHEHPVTESPMNCSVTLOLSLY
22300	32074	1	22737	ľ	323	MKSVA/GIGOTGKIEPVVLPLL
		1				WFAESGAMEGETLHTFYTOLV
		1				LMPKVMHYAQYVLLALGCVL
		ı		i		LLVPVICOIRSO/DRCWPEGSPG
		ı				*QPFAFPPA*TDIILWTEPATLM
		ı				AEVSACCAPELPPPPHRVPQIIG
		1				AQAEVNEPGGPAFRALLDPGA
22307	52675	A	22435	3	483	VRGSGNSHGEVSKGRMCLAGA
22301	12073	l^	12733	ľ	100	LPLPGACAALVLLLVILPEPLRC
		1		i		PLHTLYGGRDPVDSIRLWRQA
		l				GEFRHKSNITFNNNDTVSFLEY
	l	1	l	l		RTF\QFQPSKSHGSESDYIVMPN
		1		l		ILVLNKPRSTWISVCFVHGCIPA
	l	1	1	l		SRTGPRTESMHSKSLWKAWLA
		1	l	l		APGLPLL
		ட	L	L		AFGEFEE

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last aminu acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22308	52676	Α	22436	1	1623	RRRPRLLPGAEPCEPRVGPRRA
		l				DMGCSAKARWAAGALGVAGL
						LCAVLGAVMIVMVPSLIKQQV
	ŀ					LKNVRIDPSSLSFNMWKEIPIPF
						YLSVYFFDVMNPSEILKGEKPQ
						VRERGPYVYK/ESSRHKSNITFN
		1				NNDTVSFLEYRTFQFQPSKSHG
		l				SESDYIVMPNILVLGV\AVMME
		l				NKPMTLKLIMTL\AFTTLG\ERA
		ł				FHGTALWGEV\MWGYKDPLVN
		l				LINKYFPGMFPFKGGFGVFAEL
		1				NNSDSGLFTVFTGVQNISRIHLV
		l				DKWNGLSKVDFWHSDQCNMI
		l				NGTSGOMWPPFMTPESSLEFYS
		ı			l	PEACRSMKLMYK\ESGVFEGIP
		l				TYRFVAPKTLFANGSIYPPNEGF
		l				CPCLESGIQNVSTCRFSAPLFLS
	1	l				HPHFLNADPVLAEAVTGLHPN
	ĺ	1				QEAHSLFLDIHPVTIIPMNCSVK
		l				LQLSLYMKSVAGIGQTGKIEPV
	1	l		ŀ		VLPLLWFAESGAMEGETLHTF\
		1				YTQLVL\MPKVMH\YAQYVPS
		1		l	ł	WALG\CVLLLVPV/LSCQIRSQE
		l			1	KCYLFWSSSKKGSKDKEAIQA
		l			1	GSE*SLMTSAPKGS\VLQEAKL
22309	52677	Α	22437	198	1062	STCVEMPRNLLYSLLSSHLSPHF
	l	l		ľ		STSVTSAKVAVNGVQLHYQ\HF
		l				GEGDHAV\LLLPGMLGSGETDF
		1		l		GPQLKNLNKKLFTVVAWDP\R
		1		i		GYGHSRPPDRDFPADFFERDAK
		1		İ		DAVDLMKALKFKKVSLLGWS
		1				DGGITALIAAAKYPSYIHKM\VI
		1				WG\ANAYVT\DERPAG*YEGHP
		ı	į			RMVSQMGVEGKKESL*EALYG
	1			l	1	YDYFARTCEKWVDGIR\QFKHL
		1		I		PDGNICRHLLP\RVQ\CPPPLIVH
				l		G\EKDPLVPRFHADFIHKHVKG
		l		l		SRLHLMPEGKHNLHLRFADEF
ĺ		1				NKLA\EDFLQ
22310	52678	A	22438	223	393	AOASPPLLPP**ERSNVPCPIS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22311	52679	Α	22439	524	1516	FILEDITGYVMAPIDGLLCREPR
		1		l	\	RGLYLEGTKPMTLFQIQFLNSV
		1				GDLLDLIPSLSPTKNGSLKVFKR
		ı				WDMGHCSALIKVLPGFENILFA
		l l				HSSWYTYAAMLRIYKHWDFNII
		l				DKDTSSSRLSFSSYPGFLESLDD
						FYILSSGLILLQTTNSVFNKTLL
		i i				KQVIPETLLSWQRVRVANMMA
						DSGKRWADIFSKYNSGTYNNQ
		1				YMVLGPEESKAEPQSLTKGT/P
		l				VTLWEQIPTY/VRNISGTNLMFL
		l		1		RGKGFWA\SYNVPFHEKIYNWS
İ		1				GLSHC*FRKLGLGPTPYDFRPH
ļ		1				EPKIFRRDPR/DK*LIRHPMKYI
						MRYN\NYKE\DPYSRGDPPVNT
22312	52680	Α	22440	1	306	
22313	52681	Α	22441	698	1613	FOFFOIFPEYKNNDFYVTGESY
						AGKYVPAIAHLIHSLNPVREVKI
						NLNGIAIGDGYSDPESIIGGYAE
l						FLYQIGLLDEKOKK\YFQKOCH
				1		ECIEHIRKONWFEAFEIL\DKLL\
		ı		1		DGDLTSDPFLTSONVTGCS\NY
Į.		ŀ		1		YNFLRCTEPEDQLYYVKFLSLP
						EVRPAHPTLGN\OTFN\DGPIV*
		l				KVPCGEDTSTSPVKPW\LTEI\M
						NNYRVLIY\NG\QLGHPSVAAAP
		ŀ				DOSAPLMG\MDWKGSP\EYKK
						AEKK\VWKIFKS\DSEVA\GYI\R
		1		i		OVGDF\HQVIIRGGGHILPYDOP
		l		İ		LRAFDMINRFIYGKGWDPYVG
22314	52682	Α	22442	1	258	
22315	52683	Α	22443	366	848	
22316	52684	A	22444	1	585	ECLCVSTAVRIRHELKTMLTEL
		1	l			LRGGSFEFKDMRVPSALVTLH
	1	1	İ	i		MLLCSIPLSGRLDSDEQKIQNDI
		1		l		IDILLTFTQGVNEKLTISEETLA
			l	l		NNTWSLMLKEVLSSILKVPEGF
			l	1	1	FSGLILLSELLPLPLPMQTTQVS
		1	l	l		LPYNMHLINDCSNTF*KASDSV
			l	1		KKQKPSSPFLPKRHKITG*GGD
l	1	1		1		KTETSWSLRKYGGKNFK

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ĺ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22317	52685	A	22445	625	2700	YIRGHSTPRKLWSMHLHVQAK
			ŀ			LLQEIVRSFSGTTCQPIQHMLRR
			ŀ		ľ	ICVQLCDLASPTALLIMRTVLD
			ŀ			LIVEDLQSTSEDKEKQYTSQTT
						RLLALLDALASHKACKLAILHL
			İ	ŀ	ļ	INGTIKGDERYAEIFQDLLALVR
				İ		SPGDSVIRQQCVEYVTSILQSLC
				1		DQDIALILPSSSEGSISELEQLSN
			ŀ		}	SLPNKELMTSICDCLLATLANS
	l					ESSYNCLLTCVRTMMFLAEHD
			ŀ		1	YGLFHLKSSLRKNSSALHKLLK
				ŀ	ŀ	RVVSTFSKDTGELASSFLEFMR
				ŀ	ĺ	QILNSDTIG\CCG\DDNG\LMGSR
			ļ.		ĺ	RGAHTSRTMSINAAELNQLLQS
			l	1		TEESPEDLFLELEKLDLEHSKD
				1		DDNLYSLLD**RGLKQMLESSG
				l		DPLPLSDQDVEPVLSAPESLQN
				İ		LFNNRTAYVLADVM\DDQLKS
				l		MWFTPFQAE\EIGTDLDLVKVD
				l		LIELSEKCCSDFDLHSE\LERSFL
				1		SEPSSPGRTKTTKGFKLGKHKH
				1	!	ETFITSSGKSEYIEPAKKEFMLC
				1	i	PPPRGRGRGGFGOGIRPHDIFR\
				1		ORKONTSRPP\SMHVDDF\VAA
			İ	İ		E\SKEVVPQDGI/PSHPNRPLKVS
1			1	1		ORISSRVGFSGNRGGRGAFHSO
				1		NRFFTPPASKGNYSRREGTRGS
1				1		SWSAONTPRGNYNESRGGOSN
1		ļ		1		FNRGPLPOL\RPLKFPTG*PAPK
1		ļ		1		SLGTRAS*MVRGG\LGTFPGLS
1	ŀ			1	ì	AKLAASGGSRGKFVSGG\SGRG
22318	52686	A	22446	2487	3161	TWWLHISHIPGGKKSYSPRLGR
22510	32000	^	22770	2407	5.01	GKRFATFVLLVGGELCLTSYAP
ł				1		PSIHPILRSEAECLNYSIFSKFLA
]				1	!	PAGEFALPRSFDNCPRVRLRAL
1				1		RLALLILLTFILCWTPYYLLGM
				1		WYWFSPH\MLTEVPPSLSHILFL
1				l	l	LGLLNAPLDPSPSYGAFTLGCR
1	l			l		RGHQELSIDSSKEGSGRMLQEEI
	l		l	l		
	l		1	l	I	HAFRQLEVQKTVTSRRAGETK
22216	50607	-	22447	222	227	GISITSI*S*QSM*EQNSKSLVP
22319	52687	A	22447	232	337	
22320	52688	A	22448	I	1449	
2232 I	52689	Α	22449	1	2004	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	deletion, \=possible nucleotide insertion)
22322		A	22450	3	1558	APARDRAGGGRRALPRIPAAS ARARGITRALSRAMACRQEPQ PQGPPPAAGAVASYDYLVIGGG SGGLASARRAAELGARAAVVE SKHLGGTCVNVGCVPKKVMW NTAVIISEFMHDHADYGFPSCE GKFNWRVIKEKRDAYVSRLNA IVQNNLTKSHIEIIRGHAFFYSD PKPTIEVSGKKYTAPHILIATOG WYPSTPH-FERSRGAS.I-YISDG FFQLEEIVPAAASLVGARLPLSV EMAGILISALGSKTSLMIRHDIX URSPFDSMISTNCTEELENAGV EVLKFSQVKEVKKALISGLEAS WYTAVPGRIPVMPMIPDVDCL LWAIGGVPNTKDLSLNKLGIQT DDKGHIIVDEFQNTNVKGITAV BOVCGKALILTPVAIHAGRKLA HRLFEYKEDSKLDYNNIPTVVS SHPPIGTVGLTEDEAHKYGIEN VKTYSTSFTPMYHAVTKKKTK CVMKMVCANKEEKVVGIHM CULCOEMILOGFAVAVKMGAT
		L		-		KADFDNTVAIHPTSSEELVTLR
22323	52691	A	22451 22452	51	464	
22324 22325	52692 52693	В	22452	103	1184 148	
22323	52694	A	22454	1	3282	
22327	52695	A	22455	145	436	
22328	52696 52697	A	22456	3	3622	MPGYRAGPFEEAFDREDERQIE CQGRGVDRKITGEHITTFSGRSG PDHEITLIDSGGGFTRSRA ASEDRRKLPIIVEDEGGFTRSRA CSSPARGSRPPPSAIGCSPVAQA SDSAAGPARRTALQSLSSWLGY QIDRHSVPVYVFKSPLFSVIMAP KHKSSDAGDLDRPKRSKVLPL SEKVKVLDLIRLDKKSYAEVA KIYGKNESSIREIVKKEKEIRASF EQALHLWVEE VVELRCQDKEGDPAPEYTWFK DGIRLLENPRLGSQSTNSSYTM NTKTGTLQFNTVSKLDTGEYSC EARNSVGYPOVPGKRMQVDD
22330 22331	52698 52699	A A	22458 22459	I	20 8 321	LNISGIIAAVCGLGVCYAQRKG YFSKETSFQKSNSSSKATTMSE NDFKHTKSFII

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22332	52700	A	22460	2	1169	HTPKTEQTPIPGLEDPPLGSQLR
				l		RRPGEGETDIPSRAVPPLPPLPT
				1		LCSFPAPEVQGPPASCAPAAGT
				l		LDLLRAAGCRPGKMARRSRHR
				l		LLLLLRYLVVALGYHKAYGF
						SAPKDQQVVTAVEYQEAILAC
		ŀ		1		KTPKKTVSSRLEWKKLGRSVSF
				l		VYYQQTLQGDFKNRAEMIDFNI
		i				RIKNVTRSDAGKYRCEVSAPSE
1				l		QGQKPG\EDTVTLEVLGDVHVL
1						APAVPSCEVPSSALSGTVVELR
1				l		CQDKEGNPAPEYTWFKDGIRLL
		ĺ		l		ENPRLGSQSTNSSYTMNTKTGT
				l		LQFNTVSKLDTGEYSCEA\RNS
				l		VG*SEGPCETNASR*SQHKVGII
		l		I	1	GSRS*LWALVDFPFVGLVYSYG
		l		I	1	SEGKATFSKETSFQKSNSSSKAT
						TMSENDFKHTKSFII
22333	52701	В	22461	57	168	
22334	52702	Α	22462	1	682	
22335	52703	Α	22463	398	1465	GTFPPSGLITGKKRKALLGACD
		ĺ				CTQIVKPSGVHLKLVLRFSDFG
		l				KAMFKPMRQQRDEETPVDFFY
1						FIDFQRHNAEIAAFHLDRILDFR
1				1	1	RVPPTVGRIVNVTKEILEVTKN
		l		l	1	EILQSVFFVSPASNVCFFAKCPY
i i				l		MCKTEYAVCGKPHLLEGSLSA
İ				1		FLPSLNLAPRLSVPNPWIRSYTL
				l		AGKEEWEVNPLYCDTVKQIYP
						YNNSQRLLNVIDMAIFDFLIGN
				l	-	MDRHHYEMFTKFGDDGFLIHL
				l		DNA\RGFGRHSH\DEIPILSLLSQ
				1		CGMIKKNTLLHLQ\LLAQADYR
		l		1		LS\DVMRESLLEDQL\SPVLTEP
						HLLAL\DRRLQTILRTV\EGCIV\
						AHG\QQSVH*FDGPSGNSSGPR
22336	52704	Α	22464	2	424	GRVGVSAVIVRAWITHKYSGR
1		l		1		DWNVKWEENLLHAVAKNYTL
		l		1		LQTIPPFERPFKDHQVCLEWNM
1		l		1		GYIWNLRANRIPOCPLEN/DFRE
		l		i		DFHYNDTAGYFIIGGSRYVAGI
		l		I		EGFFGPLKYYRLRSLHPAQIFNP
				l		LLEKOLAEOIS
22337	52705	A	22465	23	323	
22338	52706	A	22466	3	287	LSVQMYAQAALDGDSQGFFNL
				ľ	Γ".	ALLIEEGT\YERCWSHSNEESFS
		ĺ		l		PCSLAWLYLHLRLLWGAILHSA
1		l		l		LIYFLGTFLLSILIAWTVQYFQS
1				l		VSGKDL
22339	52707	A	22467	1	348	.55.122
22340	52708	A	22468	1	738	
22340	122700	<u></u>	22400	L ¹	1,20	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22341	52709	Α	22469	1	507	
22342	52710	Α	22470	187	3682	SILCLLSPCVVOFGKPVVSILSS
						RSRHSPCTKKGWEGMRKHLHT
				1		ROGHK*VHV\KFP\SIMGLOGM
						DYFHPDISIFCNLAVIVRAWITH
			į .			KYRGRDWNVKWEENLLHAVA
						K\NYTLLQTIPPFERPFKDHQVC
						LEWNMGYIWNLRANRIPOCPL
						ENDVVALLGFPYASSGENTGIV
				İ		KKFPRFRNRELEATRRQRMDYP
						VFTVSLWLYLLHYCKANLCGIL
				1		YFVDSNEMYGTPSVFLTEEGYL
						HIQMHLVKGEDLA
22343	52711	A	22471	3	153	
22344	52712	Α	22472	1	408	TEPEFPGRRFRPRMAVMAP\RT
1			l			LVLLLSGALALTQTWAGSHSM
						RYFSTSVSRPGRGEPRFIAVGYV
1		1	l			DDTQFVRFDSDAASQRMEPRA
				1		PWIEQEGPEYWDEETGKVKAH
l			l			SQTDRENLRIALRYYNQSEAGS
						HTLFPR
22345	52713	Α	22473]1	823	MAVVAPRTLLLLLSGALALTQT
			İ			WAGSHSMRYFSTSVSRPGSGEP
						RFSAVGYVDDTQFVRFDSDAA
l			İ			SQRMEPRAPWMEQEEPEYWDR
			l			QTEISKTNAQIDLESLRIALRYY
						NQSED/GPPPKTHMTHHPISDHE
				1		ATLRCWALSFYPAEITLTWQRD
				1		GEDQTQDTELVETRPAGDGTF
						QKWASVVVPSGQEQRYTCHVQ HEGLPKPLTLRWEPSSQPTIPIV
				1		GILAGLVLFGAVIAGAVVAAV
				1		MWRRKSSDRKGGSYSQAASSD
						SAOGSDVSLTACKV
L	l		L	l	L	3AQU3DV3LIACKV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequenec (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		i		sequence		
22346	52714	A	22474	1	1162	NSRVDDFVAVMAPRTL\LLLLS
22340	32714	<u> </u> ^	22474	ľ	1102	GALALTOTWAGSHSMRD\FFTS
1	ŀ					VSRPG\R*EPRFIAVGYVDDTOF
					1	VRF\DSDAASQRMEPRAPWIEO
					ł	EGPEYWDQETRNMKA\HSQTD
1						RENLRIALPLTYNOSEAGS\HTL
						QMMFGCERGGSDGALSFRGLP
					1	PVTAY\DGKGFTFALNE\DLRS
	ĺ					
						WT\AADMAAQITQRKWETPPM
				1		WRS/EQEEAYLEG\QCV\DGLRR
						YL\ENGKEDAAAQRTPPKTHMT
	i			1	l	HHPIS*P*RPL*RCW\AL\AFYPC
						GDHT*PWQRDG\EDQTQ\DTEL
						V\ETRPAGDGTF\HKWAAVVVP
			ĺ			SGEESEITPANVQQ*RVLPKPLT
						LKM/WEPSSPAHPSPIVGHHLLA
						LVL/LLGAFDHLKLWVAAVML
		l				KEGTAQVGKG*KIEKGGSYSSR
						LPSQ
22347	52715	Α	22475	9	316	IPAMGLETEKADVQLFMDDDS
1		l			\	YSHHSGLEYADPEKFADSDQD
		l				RDPHRLNSHLK\ILMPFVKTCL
		l				MVLPSVQTIWKSVTDVIIAPLCT
			1			SVGRCFSSVSLQLSQD
22348	52716	A	22476	3	606	RPRTKAAMGLETEKADVQLFM
		ł				DDDSYSHHSGLEYADPEKFADS
						DQDRDPHRLNSHLKLGFE\DVI
						AEPVTT\HSFDKVW\ICSHALFEI
						\SKYRM\YRSLT\VFLGHSPWAFI
				1		CGEFSFATL\SCLAHLGF*WPFV
						KTCLM/VFLPSVQTIWEEC*QM
	İ			1		V/ISLAPLC\TSVG\RCFLFLFSLA
	l					TEPRNGILGPOVWRLGYCNTSF
						VIIT
22349	52717	Α	22477	101	432	ATROQLKAKPENLLGSISRDSG
				1		RTOEGIPARVPYPPPP/PTREPPO
						ST*PGRNGLLGCLGGGAPRPG
						WLDSEAGGAPAFHCPG*CPRRH
						RRGGTRLKOLPPPPTTVHDOPL
22350	52718	A	22478	794	1209	VKVDEVTVKHLGCORCOWSRC
	[ľ.	I	l	1	PPSCSGTCLCCGAEAPCAAOPA
		1	1	l		WPPPPAAPPPTPPAPPSD\NSSCT
			l			O*PTPRCGT*DKDKGIFTLPKGH
						P*V*SLEHGFLKLGSCP\PPPPHA
				l		LVRPPOVRFNTPHGAEPORDOL
		1	1	l		, , ,
		Ц	L	L	L	PQST

SEO ID	SEQ ID NO:	Mat	SEO ID NO:	Nucleotide	Nucleatide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		l		sequence		
		<u> </u>				
22351	52719	A	22479	3	1041	GSNCPLRQADPGPPRHTLFMDF
		1				VAGAIGGVCGVAVGYPLDTVK
		1				VRIQTEPKYTGIWHCVRDTYHR
		l				ERVWGFYRGLSLPVCTVSLVSS
		1			1	VSFGTYRHCLAHICRLRYGNPD
				l		AKPTKADITLLGCGLRTRPRVP
		l	i			DVAH*GGQSPLADADTGFLSSS
1				l		GGFRLSALLAAPPMCPVPPACP
į.					1	EPKYRGPLHCLATVAQ\EEGLC
ŀ						GLYKGSLALVLRDGHSFATYFL
		1				SYAVLCEWLSPAGHFRPDVPG
		1				VLVAGGCAGVLAWAVATPM\D
						GIKSRLEANGQGQ\NGYRGLLH
					İ	CMVTSVREEGPRVLFKGLVLN
ļ		l				CCRAFPVNMVVFVAYEAVLRL
						RPGSAHHSRSPTPRGATHQQLV
22352	52720	A	22480	38	453	
22353	52721	A	22481	287	413	DMDYRRGGWTVIHRKIGGRID
						FQRVR*DYLHGYADLLREFWL
22354	52722	Α	22482	1	393	MPKLLKWNYEGGACWGNLW
						KLPNVERLCFSEATVFRSAAQC
1					Į.	LVVGLGLLLVSRACSHEDAIEW
1						RGDSSVVNIVEDGSNAKDESKS
		1				NDTVCKEDCEESCDVKTKITRE
1			Ì		ĺ	EKHFMCIFHNVQSLGTVCGEK
1						GIDVN*NYTRRKTFHVYLPQCS
						VTGNSLRGERH
22355	52723	A	22483	505	854	
22356	52724	Α	22484	584	1009	FTQKDLFIPFEVMCDMDYRGG
1						GWTVIQKRIDGIIDFQRLWCDY
1					İ	LDGFGDLLGDAFL\FL*KEDNQ
					ł	NAMPFSTSDVDNDGCRPACLV
1						NGQSVKSCSHLHNKTGWWFN
						KCGLANLNGIHHFSGKLLATGI
						QWGTGNQKQLTWQE
22357	52725	A	22485	1	3123	MSHRARPCPAPFLFWMLGVKT
						EPAGQDRRKKARSRCCIFRVHA
1		l	ĺ			SGLWGLRFPSVEWGKPSRDKA
						QLSESPPQSVYLFAETALSAIVP
						GDGVAGGGGTGDGGPGAPYPG
						AVADMEELTIWEQHTATLSKD
			l	l		PRRGFGIAISGGRDRPGGSMVV
		l	l	Į.		SDVVPGGPAEGRLQTGDHIVM
l	1	l	l	l		VNGVSMENATSAFAIQILKTCT
	1	l				KMANITVKRPRRIHLPATKASP
1	1		l	l		SSPGRQDSDEDDGPQRVEEVD
		ŀ	1			OGRGYDGDSSSGSGRSWD
22358	52726	Α	22486	191	267	
						·

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first coden for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		097.740,217	sequence	or pepulae sequence	decision, - possible nucleotide inscrition)
		\perp				
22359	52727	Α	22487	2	609	PPPRSSEDPAKMAEVEHEEGSG
						PFRKFTLPAAVDL\DQ\RLDISY
		1				DQLEQPMQAVPVARPAAGGLN
		1				PGACGRKQ\HSPA*KRLLQGPS
		l				KGGAAPCHKPEAGEDATLRD\
		1				MII\LPEMWGS\MVGVYH\GKTF
		1				\NQVEDQGVCGRPCRLGWAGV
		1				A*CRRDQLTPSFALGLPQPEMI
		1				GHYLGEFSITYKPVKHGRPGIG
						ATHSSRFIPLK
22360	52728	Α	22488	3	461	STISLYCGLKTCPKLTCAFPVSV
1		ŀ				PDSCCRVCRGDGELSWEHSDG
1			ŀ			DFFRQPATREASHSCHRSHYDP
		l			İ	PPSRQAGGLSRFPGARSHRGAL
i		l			1	MDSQQASGTIVRFVLNNKHKH
1					İ	GQVCVSNGRTYSHGESWHPNL
		1				RAFGIVECVLCTPTPID\WPQGD
1		1				*GKCRHGCPTLHYVD*LPLEKQ
		l				VILATALTMILHQADRLEVCPA
1						FLGPEVTGELLWIPSKHQEPLC
		1				DLSSITNTSMDKCVFPMEGPIL
		l				MASPGTQTSGLLALWSVCYVL
		_				PPPSIVPQGD
22361	52729	A	22489	656	1495	TGDGELSWEHSDGDIFRQPANR
		1				EARHSYHRSHYDPPPSRQAGGL
					1	SRFPGARSHRGALMDSQQASG
						TIVQIVINNKHKHGQVCVSNGK
İ		1				TYSHGESWHPNLRAFGIVECVL
İ						CTCNVTKQECKKIHCPNRYPCK
1		1				YPQKIDGKC*QNSTCFLIPEELP
		1		i		GQSFDNKGYFCGEETMPVYES
		l				VFMEDGETTRKIALETERPPQVI
		l		1		CSTFCLA\QGILQHFHIEKISKR
		l		I		MFEELPHGHFAFVCLLGQWKIF
		l		l		TEGEAQISQMCSSRVCRTELED
00270	50000	١.	22.400		400	LVKVLYLERSEKGHC
22362	52730 52731	A_	22490 22491	1	477 294	
22364	52732	A	22491	56	4042	VFFTSMGEHLQKLIPASPACLA*
22304	32/32	A	22492	36	4042	
				1		IPQACQP*PPTPPAWQCRNLR/L AASRHOEPRGEVGGPLGHGAE
		1		1		RILAVPVRTDAQGRLVSHVVSA
	1	1	l	I		ATSRAGVRARRAAPVRTPSFPG
		1		I		GNEEEPGSHLFYNVTVFGRDLH
	i	l		1		
	l	I	l	l		LRLRPNARLVAPGATMEWQGE
	l	1	l	l		KGTTRVEPLLGSCLYVGDVAG
	l	1	l	i		LAEASSVALSNCDGLAGLIRME
	l	1	l	I		EEEFFIEPLEKGLAAQEAEQGR
	l		l	1		VHVVYRRPPTSPPLGGPQALDT
	L	Ц.				GASLDSLDSLSRAL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop eodon, /=possible nucleotide
	sequence		09/540,217	eodon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22365	52733	Α	22493	173	1148	QALGLPSLLAVLALSTTDRASF
		1		\		SAVISRSPORTVSSTRKEORPLV
		l				FAFSALRLGTNACKMDGFCSTF
		l	İ			LGAPGIRLKLLTPAGTHKSERD
		l				RTATPTRHHEKRLFCSSLSIGRQ
		1				AEEAWRADHPLPESRLFGLVPL
		1		i		GGAAWCRLAGSFCVCSLRLPRP
		1		l		IELVRAASGALLGAVRRRNHGP
	1	1				PHCFSTATLLCKAYRGGHLTIR
		1				LALG\GAPNRPFYRIVAAQNKC
		1		ŀ		PRDGPFP*SKLGPPYDP/LLPPNS
	1	1	ŀ			HGEKTPCPLN\LDRI\RHWIG\CG
		l				APPLLSPMEKLLGLAGFFP\LHP\
						MMITNAERLRRKRAREVLLAS
			ŀ			QKTDAEATDTEATET
22366	52734	A	22494	2	4985	AGATTALELPGQPSVTGVPELP
						GLPSATRALELSGQPVATGALE
			ŀ			LPGPLMAAGALEFSGQSGAAG
						ALELLGQPLATGVLELPGQPGA
						PELPGQPVATVALEISVQSVVT
					İ	TSELSTMTVSQSLEVPSTTALES
						YN/DERSMMSSYERSMMSYER
			ľ			SMMSPMAERSMMSAYERSMM
						SAYERSMMSPMAERSMMSAYE
						RSMMSAYERSMMSPMADRSM
						MSMGADRSMMSSYSAADRSM
						MSSYSAADRSMMSSYTADRSM
22367	52735	A	22495	18	174	ALIKKHFIF*IAAAKPTPPKSQV
						TLTKEFPVSSGSQHRKKEADSV
				1		YGEWVL
22368	52736	Α	22496	3	595	IKKHFIF*IAAAKPTPPKSQVTLT
1		1				KEFPVS\SGPQHRKKEADSVYG
		l				EWVPVEKNGEENKDDDNVFSS
						NLPSEVSRRNIMYFSFFIPESAH
						FIVMLCLIWDSVSLFLRAGVKR
						QGRV*DGQM\KQPAASH\LTVT
						RM/CNSLCG\TKATKWKRLRFA
						ENSVI\TSLPNIGPSLHLWEGSP
						WYSIPFSFPSLLLLKVGEVEFKR

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	l district			sequence		
22369	52737	A	22497	3	5618	SSELPGPSVTPVPQLSQELPGLP
						APSMGLEPPQEVPEPPVMAQEL
				l		PGLPLVTAAVELPEQPAVTVA
l						MELTEQPVTTTELEQPVGMTTV
	100	ĺ				EHPGHPEVTTATGLLGQPEATM
ŀ						VLELPGQPVATTALELPGQPSV
						TGVPELPGLPSATRALELSGQP
					ľ	VATGALELPGPLMAAGALEFS
						GQSGAAGALELLGQPLATGVL
					ŀ	ELPGQPGAPELPGQPVATVALE
			ļ.		ŀ	ISVQSVVTTSELSTMTVSQSLEV
				l		PSTTALESYNTVAQ
22370	52738	Α	22498	1	4685	MAPESHILASNTMETHILASNT
					l	MDSQMLASNTMDSQMLASNT
				l	ŀ	MDSQMLASSTMDSQMLATSSM
						DSQMLASNTMDSQMLASNTM
					İ	DSQMLASSTMDSQMLATSSMD
						SOMLATSSMDSQMLATSTMDS
						OMLATSSMDSOMLATSSMDSO
					İ	MLATSSMDSOMLATSSMDSOM
					[LATSTMDSQMLATSTMDSQML
						ATSSMDSOMLASGTMDSOMLA
1						SGTMDAQMLASGTMDAQMLA
					ŀ	SSTQDSAMLGSKSPDPYRLAQD
					ŀ	PYRL
22371	52739	Α	22499	291	7655	KRAPEKVDAYLYKOILLMKLP
						TKKSKKHKKHKNKKKKKKKE
						KEKKYKROPEESESKTKSHDD
1						GNIDLESDSFLKFDSEPSAVALE
l						LPTRAFGPSETNESPAVVLEPPV
l			1			VSMEVSEPHILETLKPATKTAE
1			l			LSVVSTSVISEQSEQSVAVMPEP
l			1	l		SMTKILDSFAAAPVPTTTLVLK
				l		SSEPVVTMSVEYOMKSVLKSV
l	l	l	1	l		ESTSPEPSKIMLVEPPVAKVLEP
				l		SETLVVSSETPTEVYPEPSTSTT
				l		
				L		MDFPESSAIEALR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
00000		 		0.0	laree	COVERNI DEL CAMBRELLA COMO
22372	52740	A	22500	90	2166	SQHFGRLRRVDYLRTSKMQTSS
						SRSVHLSEWQKNYFAITSGICT
						GPKADAYRAQILRIQYAWANS
						EISQVCATKLFKKYAEKYSAIID
						SDNVESGLNNYAENILTLAGSQ
						QTDSDKWQSGLSINNVFKMSS
						VQKMMQAGKKFKDSLLEPALA
						SVVIHKEATVFDLPKFSVCGSS
1						QESDSLPNSAHDRDRTQDFPES
						NRLKLLQNAQPPMVTNTARTC
						PTFSAPVGESATAKFHVTPLFG
						NVKKENHSSAKENIGLNVFLSN
						QSCFPAACENPQRKSFYGSGTI
			ŀ			DALSNPILNKACSKTEDNGPKE
					i	DSSLPTFKTAKEQLWVDQQKK
						YHQPQRASGSSYGGVKKSLGA
1		1				SRSRGILGKFVPPIPKQDGGEQN
						GGMQCKPYGAGPTEPAHPVDE
						RLKNLEPKMIELIMNEIMDHGP
						PVNWEDIAGVEFAKATIKEIVV
1						WPMLRPDIFTGLRGPPKGILLFG
			ŀ			PPGTGKTLIGKCIASQSGATFFSI
						SASSLTSKWVGEGEKMVRALF
						AVARCQQPAVIFIDEIDSLLSQR
1	1					GDGEHESSRRIKTEFLVQLDGA
1						TTSSEDRILVVGATNRPQEIDEA
1						ARRRLVKRLYIPLPEASARKQI
1						VINLMSKEQCCLSEEEIEQIVQQ
1		1				SDAFSGADMTQLCREASLGPIS
1		l				AV/SQTADIATITPDQVRPIAYID
1		l				FENAFITVRPIVSPKDL*LYENW
22373	52741	Α	22501	3	523	FFFFFPAFKMSKRG\RGWVPSG
i	1	Į.				CRKFPDFPLGLP\VGTL*SNWCL
1		l	1			TNT\GAQKTLYI\ISREGGIKGTG
		l	1			LTRLPRLPGVG\DMVMAHSSRK
1		l	l			ANPELQKKGYIPA\VVIRQRKVI
1		ĺ	I			TVGKRWRCFFIFED\NAGVIGD\
		l				NKG\EMKGFAITG\PRAKECAD
1		l	1			LWPRDCIQMLGSIAWFSQYIL
L			L	L	l	E W I KDCIQMEOSIA WISQIIE

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino seid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22374	52742	A	22502	13	946	GPSASVSFWLWSQQHSPRIRHP
22374	32/42	^	22302	13	340	RPGERRRLLHGIPHPTAEKOPA
		l				PLRPERRGGRRRRPAGRRGCA*
		l				/GGARPCSSRSS*PASTPAPRAA
		l				PPCRSPCSARPRTGPRTGAGRPR
		l				AP/HDGPPCATPDPRDOAS\PLPP
		l				RGPRPRVAPGSP\PEGHPRELPS
		1				EGSPGPRAPGORHVRPPRAVLR
		1				PRGLRGGGCPGPAHPAPRLQRR
						RAPGEAAAPGPRWPGRPGRGR
1		l				AOALVPAARSRCFWKLWEAVF
						CLPVAAAPTRRFGARORPOLW
		1				LHGHKGGOVGEVSSHSAGEEH
		1				LVPHAAPKRAREPHGRSGSPGG
		1				IASSWODGS
22375	52743	A	22503	34	823	IA33 WQDGS
22376	52744	A	22504	424	2339	LLRGTLQSYAWTSVIPDIATELR
22370	32/44	n	22304	1424	2339	DAPEKTLACMGLAIHQVTICLM
						HTFV*TFLMFAHTCTPCLTMVN
		l				VPHIHAR*GI*CIKVLLRMGH*R
					1	PFKNENAHRGTVVRVSNIKPLC
		1				TKMAFLCAACGEIQSFPLPDGK
		1				YSLPTKVIRSLTASLFILVKKAN
		1				QNTENISRQSIK*AIRLLHKRQA
		1				LDVTLLRKSIILLHDLVDSCVPG
		1				DTVTITGIVKVSNAEEGRVOLF
		1				SLFVSQ**FINVHLF*DAPPHILV
		1				VGDPGLGKSQMLQVDNQSFSV
		l				CSFACKMR**FTSEFSPLSKDSS
		1			1	SGDFALEAGALVLGDOGERPK
		1				GIISDSGTF*KAFSFSVNFLSISL
		ı				AKAGVVCSLPARTSIIAAANPV
		1				GGHYNKAKTVSENLK*VSSNIY
		1				TFNVLNHFNSIRKRNSSHLLSEH
		1				VIAIRAGKQRTISSATVARMNS
						ODSNTSVLEVVSEKPLSERLKV
						*MFLLLIQFGSV*MSKFSVWLIV
1	1	1			1	DVYPRLSTEAARVLQDFYLELR
						KQSQRLNSSPITTRQLESLIRLTE
		1	1			VCFFLWSCFFWLKGEGCALTSL
		1				LKYR*DKNFKLF*MKYOLSSHL
			1			KLGNRPNRSTAKRFISALNNVA
		1				ERTYNNIFQFHQLRQIAKELNIQ
		l				VADFENFIGSLND\QGYLLKKG
		1		1		PKVY\QLQTM
22377	52745	A	22505	73	214	I K T I V L V I W
22377	52746	A	22506	1	503	
223/6	32/40	ΙΛ.	122300	I ¹	303	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
				<u> </u>		
22379	52747	Α	22507	3	1000	ESKLEGKAAQDIKDEEPGDLGR
				ľ		PKPECEGYDPNALYCICRQPHN
						NRFMICCDRCEEWFHGDCVGIS
1			•			EARGRLLERNGEDYICPNCTIL
						QVQDETHSETADQQEAKWRPG
	ŀ					DADGTDCTSIGTIEQKSSEDQGI
						KGRIEKAANPSGKKKLKIFQPG
	İ					PGPVPTQLPVLWQVLEIAVSRSI
						SAFTLLHCISCKVIEAPGASKCI
	ŀ					GPG\CCHVAQP\DSVYC\SNDCIL
						KHAA\ATMKFLS\$GKEQKPKPK
1						EKMKMKPEKPSLPKCGAQAGI
1	ł					KISSVHKRPAPEKKETTVKKAV
						VVPARSEALGKEAACESSTP\S
1						WAS\DHNYNA\VKPEKTAAPSP
						VTVV
22380	52748	Α	22508	I	2330	MAAAEGKERAGGRASGRGRSE
	1					PPRPGARASDAGPGGLREVEGT
						GEHSVVAAGTVPRVGVGAAAR
						GEERRAGLGECGPGLGGCVAG
	}					LISGKSVQAFSVGAGRWPLLRV
						FGGNVTFCSVVSWSLCAYTLPL
						GMDDKGDPSNEEAPKAIKPTSK
	ł .			l		EFRKTWGFRRTTIAKREGAGD
	ļ					AEADPLEPPPPQQQLGLSLRRS
	1					GROPKRTERVEOFLTIARRRGR
						RSMPVSLEDSGEPTSCPATDAE
1						TASEGSVESASETRSGPQSASTA
						VKERPASSEKVKGGDDHDDTS
		1				DSDSDGLTLKELQNRLRRKREQ
						EPTERPLKGIQSRLRKKRREEGP
						AETVGSEASDTVEGVLPSKQEP
	İ					ENDQGVVSQAGKDDRESKLEG
						KAAQDIKDEEPGD\LGRPKPEC
						EGYDPNALYCICROPHNNRFMI
						CCDRCEEWFHGDCVGISEARG
		1		1		RLLERNGEDYICPNCTILQVQD
						ETHSETADQQEA*MGDLGDA\D
1		1		l		GTDF\TSIGNIRSRSLSRRPKGLK
						GRIEKAANPSGKKELKIF\OPVL
1		İ				EIAVSRSISAFTLLHCISCKVIEA
						PGASKCIGPGCCHVAQPDSVYC
						SNDCILKHAAATMKFLSSGKEO
						KPKPKEKMKMKPEKPSLPKCG
		1				AQPVQWLLLLSSWQLGTSSGL
		l				OVTLCOPPFTEHKWLKFSVLHG
						ETAGPPRAHRIRKLNKYWGAC
l						HHCMPLRPLPCLPGWPRSSPSL
	l			1		HILLWIFERPERCERO WERSSPSE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
ļ				sequence	,	
22381	52749	A	22509	149	2001	I IQPPDPRNMTLAAYKEKMKELF
22301	32143	l^	22303	177	2001	LVSLFCSCFLADPLNKSSYKYE
						AGVGDSVGGRMKASGKTVLT
		1				GRERRAQADTV\DLNWCVISD
				1		
				1		MEVIELNKCTSGQSFEVILKPPS
						FDGVPEFNASLPRRRDPSLEEIQ
						KKLEAAEERRKRGPDLVIYEIH
						SPASLPFHSSDASVLGDIPPTTY
						SNISNSNQELLLMGGSPAPEQP
						HYEANILERSETTSYDNGSDAA
						QSFKVDGRCVFYINIPIPSNEGK
l					1	VLWKAPEIATAKEPTWSPLGST
						GPRPCQILVVKSPVVHGKGPVL
						SGCGLDLLEQPTFALCFLVAGA
						VTITFLSCSGEEAREESAPKAMS
ļ						ERQARRGGAEKQGAEGRGLQA
						VTTPELPYRMGLGAAFLLKERS
ļ		1				GVLPGSFICEDFTAALKQDEEIG
						TMKGKIREAYFKSFVPTPVAFS
ĺ				1		GGVMLPTGGTLEMCGNNFGGH
	1			1		KEQAGTVLPAFAIALRVTPSAS
l		1				LVLRPLNLDQAMLQGSLGLQL
					1	ADGLFWDLASIMDWTASRIQPS
		1				VSSSMKICQAFPGRPLAILLAQC
						AYVYGTWMKLEAIILTKLTQE
						QKTKHHLFSLPCCMPADISYRQ
						RCLGEPPHCTVWNRHLESLVM
		_				MSNSTTSPCSELLIFEVPPVAPE
22382	52750	Α	22510	373	990	IQPPDPRNMTLAAYKEKMKELP
						LVSLFCSCFLADPLNKSSYKYE
						ADTVDLNWCVIFRHWKVIE\LN
				İ		KCT/SGANSFEVILKPPSFDGVP
						EFNASLPRRARPIPWEEIPVRKL
						GRRLRERREVTREADP*KQLAE
ŀ		1		1		KR\EHEERGDSKRAFRETTTFIK
						MAK/EKNLAQKMESNRENREA
		1		1		HLAAMLERLQEK\DKHAEEVR
		L_				KNKELKEEASR
22383	52751	A	22511	524	1107	LCSPKVLGLNASCQKNFLKQK
		1	1			KSIKTLSTSQ/YSVSGYSLARSE
		1			1	NAHPEFRMAMKDGELVIWDSV
		1	1		1	HPCYTVFHEQTETFSSLWSEYH
						DDFRQFLHIYSQDVACYGENLA
						YFPKGFIENMFFVSANPWVSFT
		1		I		SFDLNVANMDNFFAPVFTMGK
		1	1		1	YYTQGDKVLMPLAIQVHHAVC
						DGFHVGRMLNELQQYCDEWQ
22384	52752	Α	22512	1	1413	
22385	52753	В	22513	1	2787	
22386	52754	Α	22514	1	1638	

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence		09/540,217	coden for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22387	52755	A	22515	71	2421	YTVCYFLQAAEEDETQNQHDD
						SSSDSGTPSGPDFNYILNMSLW
					İ	SLTKEKVEELIKQRDAKGRE\V
					ľ	NDLKRK\SPSDLW\KEDLAGIC
						WRDWDKV/ENSPRNEEDVLAG
						MSGKSNLKGKVGKP\KVKKLO
						LEETMPSPYGRRIIPEITAMKAD
						ASKKLLKKK/KGVDLDTAAVK\
						VEFDEEFSGAP\VEGAGEEGIDS
						HQFPINK\GPKP\RREKKEPGS\R
						VRKTPTSSGKPSAKKVKKRNP
		1				WSDDESKSESDLEETEPVVIPR
	l	l				DSLLRRAADIHRLGGEGFDAGL
	1	1				TVFAPLFASPFPPRGAPYLKAA
		1				COTPRSGIGATAISGEALKGAV
		ł				SSGVDPRWASHLRVKLLEKSFS
		1				LQEIQSTEKVLSQQAKTLSLLG
		1				EPKLSKOLTGTCIYRKTNLALC
		l				RGPAAAKSLFOGAVRNDRNKK
		l	l			KKETSKQECTESYEMTAELDDL
		l				TEKIRKAHOETFPSLCOLGKYT
		l		l		TSLOKGMORCOILDLSSVRPIV
		ı				DVRPPELQDSKLVLVQAAKCA
		1	İ	l		PLADHCSKPVLLKEGPEDOKD
		l		ĺ		QHDVGDCWNPECQTLFDQNN
		l	ŀ	1		AAKKEESETANKNDSSKKLSVE
1		1		1		RVYHIKTOLEHILLRPDTYIGSV
1		l		1		EPLSKVAEEGPAAVGGWKGWF
		l				GTESRKDPKPTGTEEVERGVW
1		l		1		LSGREGRLLSPSVVTTMVLAEN
		ı		1		SAIRFVGGKPSSDTVPKPKRAP
		l				KOKKVVEAVNSDSDSEFGIPKK
22388	52756	A	22516	1424	4860	GRDRYGVFPLRGKILNVREASH
22300	32/36	Ι^	22310	1424	4800	KQIMENAEINNIIKIVGLQYKKS
		l		1		YDDAESLKTLRYGKIMIMTDO
1		l		1		DODGSHIKGLLINFIHHNWPSLL
		l	1	l		KHGFLEEFITPIVKASKNKQELS
		1	İ	1		
		I	1	I		FYSIPEFDEWKKHIENQKAWKI KYYKGLG\TSTAKEAKEYFA\D
		1	l	I		MERHRILFRYAGPEDDAAITLA
		1		l		
1		1		I		FSKKKIDDRKEWLTNFMEDRR
				I		QRRLHGLPEQFLYGTATKHLTY
1		1	1	I		NDFINKELILFSNSDNERSIPSLV
22222	******	ļ.	00410	0.7	200	DGFKPGQRKV
22389	52757	A	22517	87	292 903	
22390	52758	A	22518	1		
22391	52759	A	22519	3	311	

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, -possible nucleotide insertion)
				sequence	1	
00000	50750	<u>_</u>	100500		1,000	LANGE AND ADDRESS OF THE PARK OF
22392	52760	Α	22520	l .	1087	MPSAAWKRPVEGPATLRRPLGI
					1	AAASQCPSSPQPGCLIGAVNLK
					i	SSNRTPVVQEFESVELSCIITDS
						QTSDPRIEWKKIQDEQTTYVFF
					İ	DNKIQGDLAGRAEILGKTSLKI
			ľ		i	WNVTRRDSALYRCEVVARNDR
						KEIDEIAIEVT\VQVKPVTPVCR
						VPKAVPVGKMATLHCQGE*GP
			ľ		i	PPA\HYSWYRNDVPLPTDSRAN
				1	ŀ	PRFRNSSFHLNSETGTLVFTAV
					İ	HKDDSGQYYCIASNDAGSARC
						EEQEMEVCHTDKRGGFPWSW
						AAPHLWFLQDTALLPAALTGW
						CSVSVAFPGAQCKLSVDLSFWG
		ŀ				LEDGGPLLTAPLGSTPVGTLCR
						GSDPTFPFCTALADDLYESPTPA
						ANFCLDIQAFSYIL
22393	52761	Α	22521	628	1117	FFISVINGQVSSVQRLSGVGPAC
						LSCGSANPGPPPGTSPGAGAQR
						R*\PRADGSGSPQWPRGARVGG
				ļ		GRLGTGGRGRPGWROVPRRLS
						PGFGR*GGTGPGPVGTSGKRGP
						SRRRAPANDKAACWPRFPGOP
l						AS*TGFRGERGVKGFSSWGSG
						WRAWEDGGTVH
22394	52762	A	22522	2	238	
22395	52763	A	22523	131	2891	
22396	52764	A	22524	131	3103	LVSFLLFQDALPEGDASPLGPY
						LLPSGAPERGSPGKEHPEERVV
						TAPPSSSQSAEVLGELVLDGTA
						PSAHHDIPALSPLLPEEARPKHA
						LPPKKKLPSLKOVNSARKOLRP
						KATSAATVQRAGSQPASQGLD
						LLSSSTEKPGPPGDPDPIVASEE
						ASEVPLWLDRKESAVPTTPAPL
1			1		1	OISPFTSOPYVAHTLPORPEPGE
l					1	PGPDMAQEAPQEDTSPMALMD
1		l				KGENELTGSASEESQETTTSTIIT
1			1		1	TTVITTEQAP
	l	L_	l	l		ITTYTITEQAF .

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	location of first	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22397	52765	A	222525	131	2527	SASPLGPYLLPSGAPERGSPGKE HPEREXVTAPPSSSOSAEVLGE LVLDGTAPSAHDIPALSPLLPE EARPKHALPPKKKLPSLKQVNS ARKQLRPKATISAATVQRAGSQ PASQGLDLLSSSTEKPGPFGOPD PIVASEEASEVPLWLDRKESAV PITTAPLQISPFTSQPYVAHTLP QREGEGFPDMAQEAPGEDTS PMALMDKGENELTGSASEESQ ETTTSTHITTIVITEQAPGMC THOMOGORY THOM
						LLPSGAPERGSPÖKEHPEER VV TAPPSSSQSAEVLGELVLDGTA PSAHHDIPALSPLLPEEARPKHA LPPKKKLPSLKQVNSARKQLRP KATSAAT VQRAGSQPASQGLD LLSSSTEKPGPPGDPDPIVASE ASEVPLWLDKKESAVPTTPAPL USPFTSGPVYAHTLPORPEPGE PGPDMAQEAPQEDTSPMALMD KGENELTGSASESQETTTSTIIT TTVITTEQAP
22399	52767	c	22527	161	352	
22400	52768	A	22528	2	368	LGSTHASVRINILSVDVLYFIFY RRDDNKDIDSEKEAAMEAEIKA ARERAIVPLEARMKQFKDMLL ERGVRKQSLGEIFLFCISNINMS C*KVKSKVSAFSTWEKELHKIV FDPRYLLLNLI

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22401	52769	IA	22529	2	614	KDHRWESGSLLEREEKEKLFNE
22401	32709	ľ	22329	2	017	HIEALTKKKREHFROLLDETSAL
		1				TLTSTWKEVKKIIKEDPRCIKFS
		1				SSDRKKOREFEEYIRDKYITAK
		1	1			ADFRTLLKETKFITYS*FSERCE
			1		i	SESL*KMLGLAAPQTEVLCSST
ļ						
		1			i	E\RSKKLIQESDQHLKDVEKILQ
						NDKRYLVLDCVPEERRKLIVAY
		_				VDDLDRRGPPPPPTASEPTRRST
22402	52770	В	22530	201	296 1290	
22403	52771	В	22531	1	3369	OCCUPALED CONCEPTED THE
22404	52772	A	22532	3	3369	SSSVMAERGGVGGESERFNPGE
						\LRMAQQQALRFRGPAPPP\NA
1					1	AMRGPPPLMRPPPPFGMMRGPF
						PPPRPPFGRPPFDPNMPPMPPPG
						GIPPPMGPPHLQRPPFMPPPMSS
						MPPPPGMMFPPGMPPVTAPGTP
						ALPPTEEIWVENKTPDGKVYYY
					l .	NARTRESAWTKPDGVKVIQQS
1						ELTPMLAAQAQVQAQAQAQA
						QAQAQAQAQAQAQAQAQA
		1	l			QAQAQAQAQAQAQAQAQA
		1				QAQAQAQAQVQAQVQA
22405	52773	A	22533	43	347	
22406	52774	Ä	22534	1	1776	
22407	52775	Ā	22535	3	433	PLAPNHLGGGAVVLGAESHAS
		Γ.		l	1	KDVAIDMMDSRTSQQLQLIDE
		1				QDSYIQSRADTMQNIESTIVELG
		1				SIFQQLAHMVKEQEETIQR*DR
		1				VNSAHPWDPVHLLPSTOOVFIE
		ı				HLFM
22408	52776	A	22536	304	1403	SRGNATGLRTRIRVSTWVSQRH
22400	22770	ľ,	22550	150.	1	RSCPLQLLAVAAATSSPLPPPVT
l		ı				LVPPPPDTMSCRDRTOEFLSAC
		1			i	KFAARPVKNGI\QTN*GQLLRW
		ı				LVRQRSEIHPSLAKRHLGKDLS
		ı				NTFAKLEKLTILAKR\KPLFDDK
		ı				AGEIEELTYIKODINSLIKOIAO
		ı				LO\DFVRAKG\SQSGRHLQTHS
		ı			1	NTIV\VSLQSKL\ASMSNDFKSV
	1	1				
		1			1	L\EVRDRGTLKQ\QREPEERQFS
		1			1	RAPV\SALPLSPTHRAGGA\VVL
		1		1	1	GAESHASKDV\AIDMM\DSRTS
		ı		1		QQLQLIDEQDSYIQSRADTMQN
	l	1				IESTIV\ELGSIFQQLAHMVKEQ/
		ı			l	ERETIQRIDENVLGAQLDVE\AA
		1				HSEILKYFQSVTSNRWLMDKIF
22409	52777	В	22537	147	506	LILIVFFIIFVVFLA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22410	52778	A	22538	2	1397	AAPGNGLSPSSWRSELACEFOH
		l				ECTAEVRFGAVVCRGARSHSFY
					i	LEREILQ/PPGMSL/PENPVTPML
	1				l	RHLMYKIKSTGPITVAEYMKEV
	1		ł			LTNPAKGYYVYRDMLGEKGDF
	1					ITSPEISQIFGELLGIWFISEWMA
		ł				TGKSTAFQLVELGPGRGTLVGD
		1				ILRVFTQLGSVLKNCDISVHLVE
		1				VSQKLSEIQALTLTKEKVPLER
	1	l		l		NAGSPVYMKGVTKSGIPISW*S
		l				RICTD\VPKGYSFYLAHEF\LDV
		1	1			LPVHKFQKTPQGWREVFVDIDP
						QVSDKLRFVLAPSATPAEAFIQ
1						HDETRDHVEVCPDAGVIIEELS
	1		1			QRIALTGGAALVADYGHDGTK
	İ					TDTFRGFCDHKLHDVLIAPG\T
	į.	l				ADLTADVDFSYLRRMAQGKVA
	i i					SLGPIKQHTFLKNMGIDVR/L*R
	1					FF*INQMRPSVRQQLLQGYDML
						MNPKKMGERFNFFALLPHQRL
	t				i	QGGRYQRNARQSKPFASVVAG
						FSELAWQ
22411	52779	Α	22539	2	377	
22412	52780	Α	22540	2	1427	
22413	52781	Α	22541	2	568	CSSPSN/TILGVQKLNSQWRLA
		l				QDFRLINEAVIPLYPVVPNPYTL
		[LSQIPEEAESFTVLDLKDAFFCIP
		l			i	LHSDSQFLFAFEDPTDHNSQLT
		l				WMVLPQGFRDSPHLFGQAQAQ
		1				DLVHFSSPGTLVVQYMDDLLL
		l				ATSSEASCQQATLDLLNFLANQ
		1		1		GCKRVGIALGVLTQTHGTTPQP
l	1	ł	1	l	i	VAYLSKETDVAAK

SEQ ID SEQ ID NO: Met SEQ ID NO: Not	Amino acid sequence (X=Unknown,
	*-Stop codon, /-possible nucleotide
pequence 109/540,217 readon for peptide joi peptide sequence	deletion, \=possible nucleotide insertion)
sequence	
22414 52782 A 22542 I 1230	GPRGTGPKGKARSEKGCSLSHG
22414 52782 A 22542 I 1230	
	PQTNKPLVVQKGQKMEQANHP
	VGLVISVVYKDILKKIVQRETS
	HPLIHVRYAEAITGRRTAPEDK
	GSLGRDMLAKAGAIIYMNMGN
	KLPIWCHLLEEGIYLEVWALEG
	QFGRAKNACPVQIRLKDPTTFP
	YQRQYPLRPEAHKGLQDIVKH
	VKAQGLVKKCSSPCNTPILGVQ
	KPNGQWSLVQDLRLISEAVIPL
	YPVVPNPYTLLSQIPEEAEWFP
	VLDLKDAFFCIPLHYDSHDSQF
	LFAFEDPTDHTSQLIWTVLPQG
	FRDSPHLFGQALAQDLGHFSSP
	GTLVLQYVDDLLLATSSEASCQ
	QATLDLLNFLANQGYKASRSK
	AQLCLQQVKYLGLILARGTRTL
	GKERIQPILAYPHPKTLKQLWG
	FLQIT/GFCQLWIPR*SKI
22415 52783 A 22543 I 211	
22416 52784 A 22544 1 472	
22417 52785 A 22545 1 393	
22418 52786 A 22546 59 411	SWPSDKQTLVVQRGQKMEQA
	NHPDPTDHMSQLMWT/VLPQG
	FRDSPHLFGQALAQDLGHFSSP
	GTLVLQYVDDLLLATSSEASCQ
	QATLALLNFLANQGYK/LSRSK
	AQLCLQQVKYLCL
22419 52787 A 22547 43 236	TSSLLRKEDSAPSQGKSVVFTL
	TSQG*VPLWRTLQLQGPFFTPSS
	RK*LELSSANSQQHLGCPV
22420 52788 A 22548 1 1446	
22421 52789 A 22549 I 1521 22422 52790 A 22550 2 689	
	GUDDTEDD COLUMN TON
22424 52792 A 22552 636 1586	SNDRTEDDCGKHPFMSSPP\TEP
	WVCLIEGQEIDFLLDTGTTFSV
	LIPCLGRLSSRSVTIQGILGQPVT
	RYFSHLLSCNWETLLFSHAFLV
	MPESPTPLLGRDILAKAGAIISM
	KTGNKLPICCPLLEGINPEVWA
	LEGQFGRAKNAHPLQIRLKDPI
	SFPYQRQYPLRPEAHKGLQDIV
	KHLKAQDSVRKCSSPCNTPILG
	VQKLNSQWRLVQDLRLINEAVI
	PLYPVVRNPYTLLSQVPEEAEW
	FTVLDLKDAFFCIPLHSDSQFLF
	AFEDPTDHTSQLMWMVLPQRF
	RDSPHLFGQAQAQDLGHFSSPG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		Ļ.	00550	292	1.100	
22425	52793	Α	22553	292	1493	SNDRTEDDCGKHPFMSSPP\TEP
						WVCLIEGQEIDFLLDTGTTFSV
		1				LIPCLGRLSSRSVTIQGILGQPVT
1		1				RYFSHLLSCNWETLLFSHAFLV
						MPESPTPLLGRDILAKAGAIISM
			1			KTGNKLPICCPLLEGINPEVWA
		1				LEGQFGRAKNAHPLQIRLKDPI
		l	1			SFPYQRQYPLRPEAHKGLQDIV
		İ	i			KHLKAQDSVRKCSSPCNTPILG
		ŀ			\	VQKLNSQWRLVQDLRLINEAVI
		1				PLYPVVRNPYTLLSQVPEEAEW
		l				FTVLDLKDAFFCIPLHSDSQFLF
						AFEDPTDHTSQLMWMVLPQRF
1		١				RDSPHLFGQAQAQDLGHFSSPG
		l		l		TLVLQYSEIAKTLYTLIKEMER
]		1				ANTHLVEWEPEAETAFETLKQ
		1				ALVQAPALSLPTGQNFALYVIE
		1				RAGIALGVLTQTHRTTPQPVAY
22426	52794	Α	22554	2	307	
22427	52795	Α	22555	3	400	
22428	52796	Α	22556	2	78	
22429	52797	Α	22557	395	1336	SNNRTEGASSCH\PLTEPQVCLT
İ		l				IEGQEIDFLLDTGTAFSVLISCPA
		l		l		WGTGQDSVSKKNEKKRKEKA
		l				KGNHQANAEAKIAARWNPSLE
		ı		i		IPTEGPLVWNNPLQEIKPQYSLT
1		l				ETEWGLSQGHSFLPLGWLMTE
		l		İ		EVKVLIPEASQWKILKTLHQTF
		l		1		QMGIENTQQMAKSLFTGPNLL
		l				WTIRHVVKACEVCQRNNPLVH
		1				HKAPLGEQRIGHYPGEDWQLD
1		1				FTHMPKSKGFQYLLVCVDTFT
		ı	l			NWIEAFPCKIEKAQEVNKVLIH
		1	l			ETIPRFGLPQSLQSDNDPAFKA
		l				MITQGIFQGAKDTISPSLCLEATI
		1				LREGREGN
22430	52798	Α	22558	244	638	GPLNPTFLGPPCGLGGKLVFLL
		l				LRQ*VQLFRSAGSRDRCGFLDR
		1				ERNKTNQNCGKQAKKSPKDHK
		1				NPRAVGYVPFNL*QEGNLAQP
		l		1		GYMSPSPSLI*SRSRQTWGSFQ
		_	l			MILIGTKRNRPKRKSEIRERLQP
22431	52799	Α	22559	210	600	RGKRTWRERRNGWSRGFCPLP
	1	1	l			LTGSWLISRESPRLACLGEGLG
						VPGM*DLPHAWPTTLMCPELV
			1			GSWSH*LQE*SHGPSQ*VLQLL
			ŀ	l	i	RWCVWSLFLLMFGCVWSFFLL
		1	1			VSSWSRWLRSEATDLCGECYSS

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nueleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22432	52800	A	22560	2746	5211	ETSCPLVRVHYAEAITGRCTAP
						EDEGSLGOKPPARRSNNRTEGA
					i	WGKRQLMSSP/STEPRVCLTIEG
					İ	QEIDFLLDTGAAFSVLISCPGRL
						SSTSVTIQGILGQPVTRYFSHLL
	ŀ					SCNWETLLFSHAFLLMPERPTG
			l			LLGRDILAKAGAIIYMNMGNKL
1						PICCP\LLVEGINPEVWALEGOF
l	Ì					GRAKNARPLQIRLKDPTTFPYO
						ROYPLRPEAHKGLODIVKHVK
1						AQGLVKKCSSPCNTPILGVQKP
Į	į	ĺ				NGOWSLVODLRLISEAVIPLYP
						VVPNPYTLLSQIPEEAEWSWAL
]					TFKDAFFCIPLHSDSQFLFAFED
				İ		PTDHTSQLTGTVLPQGFRDSPH
						LFGQALAQDLGHFSNPGTLVLC
						YVDDLLLATSSEASCEQATPDL
						LNFLTNQGYKVSRLKAQLCLQ
	Ì		l		ļ	EVKYLGLILARGTRALSKE*IQP
	ľ				ł .	ILAYPHPKTLKOLRGFLEITGFC
١.	ł					*LWIPGYSKIARPLYTLIKETQR
	1					ANTHLVEWEPEAETTFKTLKO
l						ALVOAPALSLPTGONFSLYVRE
	l					RARIALGVLTQTHGTTPQPVAY
						LSKEIDVVAKGWPHCLRVVAA
						VAVLASEAIKIIOGKDLTV*TTY
1					l	DVNGILGAKGSLWLSDNCLLR
						YQALLLEGPVLQICTCMALNPA
						TFLPEDGEPIKHDCQQIIVQTYA
						ARDDLLEVPLTNPDLNLYTDGS
						SFVENGI/RKVSDVTILESKPLPP
						GTSAQLAELVALTWALELGKG
22433	52801	A	22561	903	4440	
22455	32801	A	22361	903	4440	DHPGHIHSSPCHERPNNVSTLPF
		ĺ				LPGEQHPILLPPRNCPGDKILEE
						NFRYNNYKRTMMSFKERLENT
			ŀ			VERCAHINGNRPRQSRGFGELL
						STAKQDLVLEEQSPSSSNSLENS
						LVKDYIHYNGDFNAKSVNGCV
		1	1			PSPSDAKSISSEDDLRNPDSPSS
			1			NELIHYRPRTFNVGDLVWGQIK
	1		1			GLTSWPGKLVREDDVHNSCQQ
						SPEEGKVEPEKLKTLTEGLEAY
						SRVRKRNRNGIKVPPRSPRSTIG
		L_				SPRPSMPSSPS
22434	52802	Α	22562	I	501	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22435	52803	A	22563	1688	2038	I SATHGLSDSCYWLVVSRAIGRN
22433	32003	^	22303	1000	2038	NLSPASPASIIHALVFLAEPOLF
						OPRILGLIDEESGHEGGRPAAKT
	1		ļ			RKLSGPWSRPAAFMSTLLINOP
	1				i	QYAWLKELGLREENEGVYNGS
						WGGRGEVITTYCPANNEPIARV
						ROA\SVADYEETVKKAREAWKI
	1	ĺ	l			
	1					WADIP\APKPKEKLVRQIG\DAL
						REEDSKLLRKPWCPLEDWGKIL
			i			LEGVGEVQEYVDICDYAVGLS
						RMIGGPILPSERSGHALIEQWNP
						VGLVGIITAFNFPVAVYGWNN
		1				AIAMICGNVCLWKGAPTTSLIS
						VAVTKIIAKVLE\DNK\LPGAICS
		1				LTCGGADIGTAMAKDERVNLL
	i					SFTGSTQVGKQVGLMVQERFG\
		l				RHYQNWRSLLELGGNNAIIAFE
	1	1				DADLSLVVPSALFACCGEQLGQ
						RCTT/SRGRLFYYMESIHDEVFK
					1	QTLKRAYGTRSRVG\NPW\DPN
	1					VLYGAISHQGRQLSMFLGAVEE
	1					AKKERLAQWVYGGKVMDRPG
	1					N\YVEPTNVTS\LDHDASIAHTE
	1					TFAP\ILYVFK\FKNEEEVFAWE*
	1					WK*NRGLSS*HSLPKDLGAESF
	1				1	RW\LGP\KGTQTCG\IVKCQPFPT
			ĺ			KWGLRLGGA\FGGEKAHLVVA
			l			RESGQ*C/WGKQYMRKVYLVL
			l			FNYS\KDLPSGPQG\IKFQ
22436	52804	В	22564	1	1434	
22437	52805	В	22565	1	1185	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22438	52806	A	22566	1	2450	GQACHASSSPLKGSRGSPPNFEI
		1				LYGPIFEDSLAPD*PKVIAGQ\IT
	1	1			1	LRDAVTGPISSPNEAGKIYOLK
		1		1		NPAVLICRVRGLHLPEKHVTW
					ŀ	RGEAIPGSLFDFALYFFHNYOA
1					1	LLAKGSGPYFYLPKTQSWQEA
					İ	AWWSEVFSYAEDRFNLPRGTIK
		ı				ATLLIETLPAVFOMDEILHALR
					1	DHIVGLNCGRWDYIFSYIKTLK
						NYPDRVLPDRQAVTMDKPFLN
						AYSRLLIKTCHKRGAFAMGGM
i		ı				AAFIPSKDEEHNNQVLNKVKA
						DKSLEANNGHDGTWIAHPGLA
		ŀ				DTAMAVFNDILGSRKNQLEVM
						REQHAPITADQLLAPCDGERTE
1						RKFNALMESEKGQSQNLWKFA
						VYSGLRHGELAALAWEDVDLE
				1		KGIVNVRRNLTILDMFGPPKTN
				1		AGIRTYTLLQPALEALKEQYKL
1					ŀ	TGHHRKSEITFYHREYGRTEKQ
i					ļ.	KLHFVFMPRVCNGKQKPYYSV
		1			l	SSLGARWNAAVKRAGIRRRNP
		1		1		YHTRHTFACWLLTAGANPAFIA
						SQMGHETAQMVYEIYGMWID
						DMSDEQIAMLNARVMVYDVA
1					l	SGNALFISELGPLPENVTWLSPE
				1	l	GEFOKWNGTAWVKDTEAEKL
				1	l .	FRIREAEETKNNLMQVASEHIA
	ł	i				PLQDAADLEIATEEEISLLEAW
						KKYRSIPLCSKVVTMMFDDRP
l	ŀ					GCPVGFPMWGRGGFDRMPPG
		1		1		WGGRPMPPSRRDMSPHRGPPP
22439	52807	A	22567	1	941	MIPNVOGPMRGLEGPNYIFIKIS
22439	32807	l ^A	22307	l'	941	KNRRYTMAVTSVGLASSLCRV
1					1	VRPCRAPAGVVSRGSPDDEWL
1					l	
ĺ						FCPQARIQQQQDID/NGTLPDFIS ETASIRDA/DWKIRGIPADLEDR
						RV/EITGPVERKMVINALNAI\F
	I					
	1		l		į .	MADFEDSLAPDW/NKVIDGQIN LRDAVNG/TISYTNEAGKIYQL
					l	
1		1				KPN/PA VLICRVRGLHLPEKH/V
1		1				TWRGEAIPGSLFDLA/LYFFHNY
1	1	1	1	1	I	QALLAKGSGP/YFYLPKTQSWQ
1		1				EAAWWS/EVFSYAEDRFNLPRO
1	1	1	1		1	T/IKATLLD*TLAGGFQMVDPSA
				1		LGG*TGPWDHLRYIKVENIPKV GEEGDAETTGGTAV
22440	52808	A	22568	2	393	GERODAETTOGTAV
22441	52809	Â	22569	1066	1824	
22442	52810	A	22570	2	165	
22443	52811	A	22571	3	902	
22773	102011	٠,,	1		P	

SEQ ID NO:	SEQ ID NO: of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22444	52812	A	22572	1	2034	
22445	52813	A	22573	176	873	RLFGEMSKLN/HNLSEVMSKLE
	1					KQHSSKVFVVKRLSSSSRRSLVI
	1					SPPVRTSTVSSPLTSPTSPSTLSS
	i				1	KSKSESVSATEDLAPDAAQGED
	1					NSEIKELLEDEEIGKEGSEASSS
	1					EEEEPLLTCNGPSQAQPSPIIESA
		ŀ				RSQEEVPPSSPALSPGGALSPSG
	1				l	HPLSSAAEVVLRTRTAVPQAKG
i	1					NNVVITSYMTNRGFFEDKKATF
	İ					APSFLMNIKGNKTSVVKNSILE
						QGQLTVN
22446	52814	A	22574	1	1890	
22447	52815	A	22575	2	1279	LSVFPRVRNFTTQAECH/KLFK
	1				ļ.	VLKNFLIAGGGVHESSKRAS*T
						LQFIYY\SEWYGHEELKAIVWN
						NDLLWEDYEEKLADQAVRTM
						EIYVAQFSEIKERIAKRGRKLVD
						YDSARHHLEAVQNATKKDEAK
		İ				TAKAEEEFNKAQTVFEDLNQEL
	1					LEELPILYNSRIGCYVTIFQNISN
	ŀ				1	LRDVFYREMSKLNHNLYEVMS
		l				KLEKQHSNKVFVVKGLSSPSTL
		1				SLKSESESVSATEDLAPDAAQG
		l				EDNSEIKELLEEEEIEKEGSEASS
		1				SEEDEPLPACNGPAQAQPSPTT
	l	l				ERAKSQEEVLPSSTTPSPGGALS
1					1	PSGQPSSSATEVVLRTRTASEGS
		1				EQPKKRASIQRTSAPPKPPEKPV
1	l	1			l	RTPEAKENENIHNQNPEELCTSP
		1				TLMTSQVASEPGEAKKMEDKE
	l	1			<u> </u>	KDNKLISANSSEKYIFQRPSVLI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop eodon, /=possible nucleotide
	sequence		09/540,217	eodon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22448	52816	A	22576	1	1791	MPGARTSSSGASENHRARGOG
						GGPQGVGRMAEGKAGGAAGL
		l				FAKQVOKKFSRAQEKAL*KLG
	١.	l			1	KAVETKDERFEQSASNFYQQQ
		ļ	l			AEGHKLYKDLKNFLSAVKVMF
		1				ESSKRVSETLQEIYSSEWDGHE
		i				ELKAIVWNNDLLWEDYEEKL\
						ADQAVRTMEIYVAQFSEIKE\RI
						AKRGRKLVDYDSARHHLEAVO
		l				NAKKKDEAKTAKAEEEFNKAO
		1				TVFEDLNQELLEELPILYNSRIG
						CYVTIFONISNLRDVFYREMSK
			İ			LNHNLYEVMSKLEKOHSNKVF
			İ			VVKGLSSSSRRSLVISPPVRTAT
		l		1		VSSPLTSPTSPSTLSLKSESESVS
						ATEDLAPDAAQGEDNSEIKELL
						EEEEIEKEGSEASSSEEDEPLPA
						CNGPAQAQPSPTTERAK\SQEE
		l				VLPSSTTPSPGGALSPSGQLSSS
						ATEVVLRTRTASEGSE\RPKKT
						ASIQRTSAPPRRPPPPRATASPRP
						CSGNIPSSPTASGGGSA\TSPRAS
						LGTGTASPRTSLEI*PNPE\PPEK
						PVRTPEAKENENIHNQNPEELC
			ľ			TSPTLMTSQVASEPGEAKKME
			ł	İ		DKEKDNKLISADSSEGQDQLQV
						SMVPENNNLTAPEPQEEVSTSE
22449	52817	Α	22577	2216	2568	RPSLGLGM/C/TIWQ/TSLA/GKH
						RRTIETHTDDYRISLEQGTPGLE
			1			PPLNALTAISKKIGCNGFFPFQIR
						PGKNETDGRMFSPAIGIVEDPV
						TGNANGPMGAWLVHHNVLPH
						DGNVLRVKGH
22450	52818	Α	22578	108	532	WGSKGNLLYSHQTHSVSHQFS
						LVACCLELEYSPFSFPYACLLSS
						LMRLEKHQELNLWPAHQRSMR
						HPQDTLVVFS/HHGRVLL*DKT
	l					SRGTIRQQHSRFTKICCSAATAA
						VTQANRVWSGPLANSNRPAAE
						GPVLAAFMGWL
22451	52819	A	22579	1	1125	

SEQ ID	SEQ ID NO:					Amino acid sequence (X=Unknown,
NO:	of peptide	hed	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22452	52820	A	22580	1	985	MCARGSLWGCYSGLERKCEAA
		l				LPDRVVSSKSACGAVSQDLLQ
		1				AOGRRAGPPSWLYADPVHTGT
						IECGGHWCLNAALLDFQSPQRL
		ŀ				LTYLHCGLQRPMESDPPELAPM
						VHHQEGIYLNIRKAMYDKPTA
					i .	NKLNWDKSNVSSLITGTRQEET
		1				TANGTTFATIHLQFFIQIPAVPR
ŀ						KTLLNRADYYFPEHNPSLTLVI
						NRRITKFSEPNWEAPPSRGRLTP
		1				HTAGYSSETKIPEERSDSSIRGS
		l	l			RKSAVLQPLLLDSQANRVWSG
		1				PLANSNRPAAEGPVC*LWWVT
						LHGPLKATMQVCQQPLW
22453	52821	Α	22581	294	934	YPPSPAELTVCRLRQLQLRDVL
		l				YCIAYPSGPGVKPFRGNSA/G/V
						VFPADNLSEAQMQLI/ARELGH
1		l				SETAFLLHSD/DSDVRI/RFTPTV
		l				EVPI/CGHATVAAHYVRAKVL/
		l				GLGNCTIWQTSLAGKHR/VTIE
		l				KHNDDYRISLNALTAIS/KKIGC
		l				NGFFPFQIRPG/KNETDGRMFSP
		l				AIGIVE/DPVTGNANGPMGAWL
22454	52822	В	22582	1163	5585	VH/HNVLPHDGNVLRVRQ*GPR
22454	52822	A	22582	2	429	AACAAAMSLVIPEKFOHILRVL
22433	32023	<u>ا</u> ا	22363	1	427	NTNIDGRRKIAFAITASSSSSSS
ŀ		1				SSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSSS
l		1				SSSSSSSSVNLLSNGLYLVKG
		1				VGRRYAHVVLRKADIDLTKRA
		1			i	GELTEDEVERVITIMONPROYKI
		1	,		l .	PDWFL
22456	52824	A	22584	84	424	GVHAASDKPCWITTSQTPDVQ
		l				HRW/WLLISSML*TAPNRRSDK
		1			l .	AFTPHPTNHAGSQLRKHPTCSI
		1			l	DGVIDIFHAVNTFVQHHMEETL
		l				KAFHVTGHHILIARHGLGVGKE
		l				NTEHAANVVNDQRNSGFFCRF
		1				QQTFCQTCCQLIQRFVNARLRD
		l				FISC
22457	52825	Α	22585	88	381	IQVLANGLDNKLREDLERLKKI
		l			\	RAHRGLRHFWG*VGGLISLPTS
		l				TQHSSYSLFFFPNLLFLLCMTCD
1						SSLFTCSLRVRGQHTKTTGRRG
		L				RTVGVSKKK
22458	52826	Α	22586	355	699	FIDTTEAGSHFLYALVRFAATIT
						TITDDGARRLRTLFQRQQADLV
		1	[SVSKSGFFTADCTHTNALIDVV
		1		1		RTIFNNAVF*NPGFVIARLKIEIA
			į.			IIKATFGQLTEDREQVLMFQAV
l		_				RR

SEQ ID NO:	of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide scquence	deletion, \=possible nucleotide insertion)
22459	52827	Α	22587	1389	1532	SVLPGRPDARQYRQTRQSW*R WSLRLRGSYPVAGRSLSRRHRQ TERF
22460	52828	Α	22588	1	427	LVHQQWIQTKKKSLIYPFCHSP SQRVVEVAALDQWLAMLYPR VEHQAVLAVVVVVVGPLVVRP YHTRRDAVRLLRQPAVFIVAV DIVRILRDAVIRPHCECTVRVIR SGSGSRLPAGSVAYGMRMSRS GARVRTLSILPVAL
22461	52829	Α	22589	I	915	
22462	52830	Α	22590	1166	1321	
22463	52831	Α	22591	813	977	
22464	52832	Α	22592	3485	4237	
22465	52833	A	22593		499	ARAAACAAAMSLVIPEKFÖHLL RIVLNTNIÖGRRKIAFAITAIKG VGRRYAHVVLRKADIDLTKRA GELTEDEVERVITIMONPTQYK RIPDWFLNROKDVKDGKFQFGS *ANGLSTTKLIKENLKPLKEDSG PIKGLRQLPWGLRVRGQHTKT TGRRGRTVGVSKKK
22466	52834	Α	22594	1	1197	
22467	52835	Α	22595	I	1905	
22468	52836	A	22596	37	514	WTWWLRCYENRHRHCHLLQQ LPTSPHQHAVATPTTVAVPTAP AATPPVVVAVVVAPVVVVVVP VVPVVASVPFVEPGSDVIVCAE MDEQWGYVGAKSRQRWLFYA YDSLRKTVVAHVFGERTFDVVI WMTDGWPLYESRPAFTGRDC VDDAFFAVAGV
22469	52837	Α	22597	1	471	
22470	52838	Α	22598	2	323	SRIQPGSDVIVCAEMDEQWGY VGAKSRQRWLFYAYDSLRKTV VAHV/FGERTIRTLTRRLFRCLR SSHDLVSSPVSHRRRRAGSRQR HRRARGGENVEYQWGALRHG
22471	52839	Α	22599	2033	2310	ARYVGNAGCTGGALFGRSQTI REAVG/ERRLAESRDVAALDTP CDPEIKMAEWMQTLKETGSTS GHIVTRRISVRISHVTPGPASQD GPDVQQR
22472	52840	В	22600	I	4891	
22473	52841	В	22601	7	1233	
22474	52842	Α	22602	193	492	
22475	52843	Α	22603	1	486	L

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22476	52844	A	22604	62	443	RSDRLLIHGFSFPAMGFRLRKT VVAHVFGERTMATLGRLMSLL SPFDVVIWMTDGWPLYESRLK GKLHVISKRYNTORIERHNLNLR OHLARLGRKSLSFSKSGTACCG
22477	52845	A	22605	971	1444	HCLTROIPMLHHQGNQLF SMGSGDMRSSPSRAAALRQLPR PFRQLAVRMEMISRFRSAKEQT KSLRKWRKAKSIF*SVAQTGFN VLVCTTIIETGIDSRQPTLLSLNA RITSVCAAAPVTRSRTFASSGY AWLLTPHPKAMTTDAKNVLKQ
22478	52846	A	22606	1	1047	LPRWKISGQVRAGTARSGDSRR W
22479	52847	A	22607	318	598	LTGFRRLCDLRFGGGAVRSVLA RSREFILQH*FRRIVTGSMGRSL VPSAPLR/PSIVLVLFRKAW*LV KTQRKMHVVSIVQKKASCWYP LLGCL
22480	52848	Α	22608	1	2652	
22481	52849	Α	22609	1	852	
22482	52850	Α	22610	1	807	
22483	52851	A	22611	1	1740	
22484	52852	A	22612	I	3144	

SEO ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22485	52853	Α	22613	173	1983	PVQNGSSPLGALPGAPLIFGPSV
						PAISRPGSCSPEWPTLDTSGYSA
						PR/WLFTAYAGSEAVFAHVFGD
						ALGDAGLLMSLLSPFDVVIWM
				l		TDGWPLYESRLKGKLHVISKRY
						TORIERHNLNLROHLARLGRKS
1				Į.		LSFSKSVEOHDKLESLPGOIDKS
						LDEASLSLRAGSLRTITHILLPLL
						RPAILSALIYSFVRAITTVSAIVF
						LVTPDTRVATAYILNRVEDGEY
		ļ		i		GVAIAYGSILIVVMLAIIFIFDWL
						IGESLLSRFEKTVDDHSQSRLPA
1						EGSILSQDISGVKLFSLLTPARG
1						YVPIRAPCPFLCASYRFFCVARQ
1			ŀ	ŀ		QGLDAALIDAPLAVDALTPLSE
				i	l	LLSHVGQAPCSVPVVDEDQQY
				l		VGIISKGMLLRALDREGGVRVP
			ĺ		1	VDYILNGFHPLLLGMPAPLAIIV
		l				FALIACRFPGSNGRIRRSRRIRH
		l				LHKMPDATLSASYQAYVRHQT
1						NVTALVLRSSHRFRKDRHCWG
1	ļ.	l				FVFLQQPILRATGQFEIQTFRYL
1				1		SVFSVQRHKTNGKTALMSGNV
						ADKAVNRATGPTACVWLFSLR
	l	ı	l			NSSMVTTRFCGLCEWADOOGA
	İ				1	RPARISENGKRGYSYLYDVHM
)				GAQSGETRGRQGEYASKPPSRE
22.406	52054	l .	22614		2149	GAQSGETRGRQGETASKFFSRE
22486	52854	В	22614	I.		
22487	52855	A	22615	1	1815	
22488	52856	Α	22616	2014	3411	E A LA COLLO DEL LA COLLO DE L
22489	52857	Α	22617	1	922	MVICQPINGGWQWPPSTGTDP
	l	1	i			APYFRIFNPTTQGEKFDHEGEFI
	l		1		1	RQWLPELRDVPGKVVHEPWK
1		ı			1	WAQKAAAPPQTRLHSTPTTIPH
1		l			i	RVPGHTPNLKPIWEVDLPKFSH
		1	1		1	KATSHATDKPSDTTQNDQIVIA
1		i i			1	YMEPPEIVHETTTSSTHPSLYIIL
		l				KRQTRELRPQSVTSRIQPGSDVI
1		l				VCAEMDEQWGYVGAKSRQRW
		l	1			LFYAYDSLRKTVVAHVFGERT
		l		l		MATLGRLMSLLSPFDVVIWMT
		İ				DGWPLYESRLKGKLHVISKRYT
1		1				ORIERHNLNLROHLARLGRKSL
1	l	ı		1		SFSKSVELHDKVIGH\YLNIKHY
L	l					DI DIONO VELLIDIA VIOTA LENIANT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22490	52858	Α	22618	1458	2185	YGHEWRWMPGNRPH\YGRWP
		l			i .	QHDFPPFKKLRPQSVTSRIQPGS
		l			i	DVIVCAEMDEQWGYVGAKSR
	İ	l		l		QHWLFYAYDRLRKTVVAHVF\
	ŀ	1				GERTMATLGRLMSLLSPFDVVI
	ľ	l		l		WMTDGWPLYESRLKGKLHVIS
		l		l		KRYTORIERHNLNLROHLARLG
		l				RKSLSFSKSVEKITSGPTTNATP
		l		1		GTOKTATDKRGGKKLPIQIPIGP
		l		ļ		ISRIASGAAINTISIGLRKFLVTA
		ı				GVMLSTQRSIYDRHQVITSAGIT
22491	52859	A	22619	493	1518	
22492	52860	Α	22620	963	1597	
22493	52861	Α	22621	1	3052	MATDSVGLNCTPAPHARDVLL
1		l				WRPSSLPVSTSCVFPLGRIGDVS
	1	ı		ì		DESDDKTEAPTPHRLEKAREEG
		ı				QIPRSRELTSLLILLVGVSVIWF
		1	İ			GGVSLARRLSGMLSAGLHFDH
		1				SIINDPNLILGQIILLIREAMLAL
		1			l	LPLISGVVLVALISPVMLGGLVF
		1				SGKSLQPKFSKLNPLPGIKRMFS
		l				AQTGAELLKAILKTILVGSVTG
		l			ĺ	FFLWHHWPQMMRLMAESPITA
					l	MGNAMDLVGLCALLVVLGVIP
						MVGFDVFFQIF
22494	52862	Α	22622	1	1233	MANDLPHSNSGDGQKPSSPDLF
		ŀ		1		TAPLLADPTVGFLPNDAEELFIF
				l	İ	LTEITEITIPCRVTDPQLVVTLHE
		l		İ		KKGDVALPVPYDHQRGFSGIFE
				!		DRSYICKTTIGDREVDSDAYYV
				l		YRLQAVPPVQLLTGWCVTAKA
		l		į.		PPDISAISALTAVKHGNCSSLTP
1		l				LLNPPGSDVIVCAEMDEQWGY
1		l		ŀ		VGAKSRQRWLFYAY\DSLRAR
i		1				RVVAHVFG*RTMATLGRLMSL
		l		Ì		LSPFDVVIWMTDGWPLYESRL
		1	1			KGKLHVISKRYTQRI\ERHNLNL
		l				RQHLARLGRKSLSFSKSVELHD
					l	KPSPQHREGVKDVMRFNEPMY
		1	ĺ			PLESRPRLYVCHLRNGAQASGR
		l	1			RCQLPMGCACHPVSQRNDRPN
		l			l	RRPERHATRKTTTRQRQQNSRQ
	1	1		l		HKQTQKKAKPNKRRATERGKG
	L	<u></u>	l		l	PQQKAERPKRHTHCFLKLQ

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nuclcotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
l	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22495	52863	A	22623	3	1386	HEASSPLAMASLTVKAYLLGK
22493	32803	^	22023	ľ	1360	EDAAREIRRFSFCCSPEPEAEAE
						AAAGPGPCERLLSRVAALFPAL
						RPGGFQAHYRDEDGDLVAFSS
l						
l					Į.	DEELTMAMSYVKDDIFRI*IKE
						KKECRRDHRPP\CAQEAPRNMV
						HPNVICDGCNGPVVGTRYKCS
l						VCPYY\DLCSVCEGKGLHRGHT
						KLAFP\SPFGHL\SEGF\SHSRWL\
					'	RKV\KHGHF\GWPGW\EMGPPG\
					1	NWEPTFLPRAG\EARPGP\TAES
ı						ASGPSEDPSVNFLK\NVGESVA\
			0			AAFSPL\NFEVDIDVEHGGKRSR
			0			LTPVSPESSSTEEKS\SSQPSSCC
			١.			SDPSKPGGNVEGATQSLAEQM
						RKIALESEGRPEEQMESDNCSG
				l	i	GDDDWTHLSSKEVDPSTGELQ
						SLOMPESEGPSSLDPSQGGNPQ
				ĺ	i	GWKEAALYPHLP\PEVEPAA*F
						ESLSOMV VPWGFS*LKGGWVH
						QGSCRPRNYDI\GAAL\DTIPVIQ
22496	52864	Α	22624	1032	1127	FYAKAEAGDRTVA*PHKPKMF
						LSIHHRVAKI
22497	52865	A	22625	121	561	VFPGPLGSLVDICGPSRVPFPWV
						VPAVLPVOFVORLPRFFRCPPV
					1	LVTSVIRSISTLIQALGSSPFARR
l					ĺ	YWGNLG*FLFLGVLRCFSSTGS
			-			PH*PMNSVNDSVSKHT/EVTPF
l			Ì			GN/PGNNGSYHLTDAYRRLARP
						SSPLT\PGHPPLYG
22498	52866	Α	22626	243	454	ATAGCPSCSATEGVVRNGKST
12450	12000	ľ	22020		[AGHORYLCSPCRKTWOLOFTY
l					ł	TASQP/GARIMGVGLNTVLRHL
	1		1		1	KNSGRSR
22499	52867	Α	22627	3	640	
22500		_	22628	236	410	
		-		116		
	52868 52869	A A				

NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
222502	52870	A	22630	276	1905	IFI YSSHQYGCPSCSATDGVV RNGKSTAGHQRYLCSHCRKTW QLQFTYTASQPAGQ*RTAIATVI CCNNCQRRHITNSRCHTDNCRR ADSASSHATSGSSRGCAGSGC CGASSTRRCLGPVCCFRGSKRS WRRQHEGHIKDCRKLVFHNFSFI HRMDNTAVPPVQLLTGWCYTA KAPPDISAISALTAVKEGNCSSL TPLLNPPGSDVICAEMDEQW GYYGAKSRQRWLFYAYDSLEK KAPPDISAISALTAVKEGNCSSL LSPFDVVIWMTDGWPLYSERL KGKLHVISKRYTQRIERHNLNL RQHLARLGRKSLSFSKSVEH KGKLHVISKRYTQRIERHNLNL RQHLARLGRADALDAPLAVDAQTP LSELLSHYGQAPCAVPVVGMEL ESELLSHYGQAPCAVPVVGMEL GYVGIISKGMLLRALDREGGV RYPVIYPKRRRTITDVCMPLSIG RVPVIYPKRRFRITDVCMPLSIG EKELKIPTPEPDFERRSVYJCE HNTNGAMGIIVNKPLENLKIEGI LEKLKITPEPDESIRLDKPVML GGPLAEDRGFILHTPASNFASSI RISDNTYMTTSRDVLETLGTDK GYBDVLVALGYASWEKGQLEQ EILDNAWLTAPADLNIEKTPIA RYBCARLGVOLTHPAQD GTPDWWIIERLLKEWOPDEIIVG LPLNMDGTEQPLTARARKFAN RHGRFGVEVKLHDERLSTVEA RSGLIEEGGGYRALNKGKVDSA SAVIILESYFEGGY

NO: of peptide sequence hod in USSN (09540,217 location of first period sequence) in USSN (09540,217 location of preptide sequence) for peptide sequence period deletion, =possible nucleotide in peptide sequence period deletion, =possible nucleotide in peptide sequence period deletion, =possible nucleotide in period deletio	wn,
22503 52871 A 22631 399 2337 PEVQGLQQAGAAARPC APRVVCEGRLRGSGLQS ASV*ATDGVVRGKSTA YLCSHCRKTWQLQFTYY AGG*REKEEKEEKEEKE KRRREGEKEEKEEKE FKKKKKKKKKKKKKKKKKKKK KKKKKKQLSIVTMS WRNEARTTGACLPSPTY PIVSTTYTAFAAATLAK RASSKQVPLLVVLWVV WFVKAGSVALGSSPGCV CKLLTGWCYTAKAPPDI LTAVKHGNCSSLTPLLNI VIVCAEMDEQWGYVGA RWLFYAYDSLRKTVVAH RTMATLGRLMSLLSPFD MTDGWPLYESRLKGKLI KARSKQQKDADTAPPQI TPGTYHAARDTRQASPR QHRQADGDKRQTRHAE RTHTGRRKTIKETATQQ, QTRRTARTHRRTARHTA RTEVAARQTTEKDNEGS TQKTPSATATAHTPRTT VQARHTSAREENHTRQH RETTTAMTNTPHYHAE SQQQTTSTERPTIDSSTT. HTPRTVROERRMTQHEK STTTNRTTTTGATQAYQHTQI	
22503 52871 A 22631 399 2337 PEVQGLQQAGAAARRPC APRVVCEGRLRGSGLQS ASV*ATDGVVRNGKSTA YLCSHCRKTWQLQFTY1 AGQ*REKEEKEEKEEKEE FKKKKKKKKKKKKKKKKKKKKKKKK	ertion)
APRVVCEGRLRGSGLQS ASV*ATDGVVRNGKSTA YLCSHCRKTWQLQFTY1 AGQ*REKEEKEEVEEEEE KRREGEKEEKEEKEE KRREGEKEEKEEKEE FKKKKKKKKKKKKKKK KKKKKKKKKK	
APRVVCEGRLRGSGLQS ASV*ATDGVVRNGKSTA YLCSHCRKTWQLQFTY1 AGQ*REKEEKEEVEEEEE KRREGEKEEKEEKEE KRREGEKEEKEEKEE FKKKKKKKKKKKKKKK KKKKKKKKKK	VGLA
ASV*ATDGVVRNGKSTÄ YLCSHCRKTWQLQFTYT AGQ*REKEEKEVEEEEI KRRREGEKEEKEEKEEKEE EKKKKKKKKKKKKKKKK KKKKKKKK	
YLCSHCRKTWQLQFTYY AGQ*REKEEKEEVEEEEE KRRRGGEKEEKEEKEEKEE FKKKKKKKKKKKKKKKKKKKKKKK	
AGO*REKEKEKEVEEEEE KRREGEKEKEKEKEKE KRKKKKKKKKKKKK KKKKKKKKKK	
KRREGGKEEKEKEEKEKE FKKKKKKKKKKKKKKKKKKKKKKK	
FKKKKKKKKKKKKKKK KKKKKKKKKKKKKKKKKK KKKK	
RKKKKK KQLSLVTVMS WRNEARTTGACLPSPTV PIVSTTVTAFAAATLAK RASSKQVPLLVVLVWVV WFVKAGSVALGSPGCV CKLLTGWCVTAKAPDDI LTAVKHGNCSSLTPLLNI VIVCAEMDEQWGYVGA RWLFYAYDSLRKTVVAI RTMATLGRLMSLSPFD MTDGWPLYESRLKGKLI KARSKQOKDADTAPPIQ TPCTYHAARDTRQASPR QHRQADDDKRQTRHAE RTHTGRKTHKETATQQ, QTRRTARTHRRTARHTA RTEVAARQTTEKDNEQSS TQKTPSATTAHTPRRTI VQARHTSARERNHTRQF RETTTNAHTHTPHYHAE SQQQTTSTERPTIDSSTT. HTTRTVROERRMTQHKFI STTTNRTTTTGAXYQHTQ	
WRNEARTGACLPSPTY PIVSTTVTAFAAATLAK RASSKQVPLLVVLVWVV WFVKAGSVALGSSPGCV CKLLTGWCYTAKAPPDI LTAVKIGNCSSLTPLLNI VIVCAEMBEQWGYVGA RWLFYAYDSLRKTVVAI RTMATLGRLMSLLSPFD MTDOWPLYESRLGKLI KARSKQOKDADTAPPIQ TPOTYHAARDTRAASPP QHRQADDDKRQTKHAE RTHITGRRKTIKETATQQ, QTRRTARTHRRTARHTA RTEVAARQTEKDNEGS TQKTFSATATAHTPRTI VQARHTSARERNHTRQI RETTINAHTNIPHYHAE SQQQTISTERPTIDSSTT HTERTVROERRMOHEK STTTNRTTTTGAYQHTQI	
PIVSTTVTAFAAATLAKC RASSKQVPLLVVLVWV WFVKAGSVALGSSPGCV CKLLTGWCVTAKAPPDI LTAVKHGNCSSLTPLLNI VIVCAEMDEQWGYVGA RWLFYAYDSLRKTVVAI RTMATLGRLMSLLSPFD MTDGWPLYESRLKGKLL KARSKQKDADTAPPIQ TPQTVHAARDTRQASPR QHRQADGDKRQTRHAE. RTHTGRRKTIKETATOQ. QTRRTARTHRRTARHTA RTRVAARQTTEKDNEQS TQKTPSATTATHTPRITT VQARHTSARERNHTRQH RETTTNAHTHTPHYHAE SQQQTTSTERPTIDSSTT. HTTRTVRDERRMTQHKH	
RASSKOVPLLVVLVWVV WFVKAGSVALGSSPGCV CKLLTGWCVTAKAPPDI LTAVKHGNCSSLTPLLNI VIVCAEMDEQWGYVGA RWLFYAYDSLRKTVVAH RTMATLGRLMSLLSPFD MTDGWPLYESRLKGKLI KARSKOQKDADTAPPIQ TPOTYHAARDTROASPR QHRQADDDKRQTRHAE RTHTGRRKTIKETATQQ, QTRRTARTHRRTARHTA RTRVAARQTTEKDNEQS TQKTPSATTATHTPRTI VQARHTSARERNHTRQL RETITINAHTNIPHYHAE SQQQTTSTERPTIDSSTT. HTPRTVROERRMOHEKI STTTNRTTTTGAYQHTQI	
WFVKAGSVALGSSPGCV CKLLITGWCYTAKAPPDI LTAVKHGNCSSLTPLLNI VIVCAEMDEQWGYVGA RWLFYAYDSLRKTVVAI RTIMATLGRLMSLLSPFD MTDGWPLYESRLKGKLL KARSKQKDADTAPPIQ TPOTVHAARDTRQASPR QHRQADDKRQTRHAE. RTHTGRKTIKEFATAQQ, QTRATARTHRRTARHTA RTRVAARQTTEKDNEQS TQKTPSATATAHTPRRTT VQARHTSARERNHTRQH REITTIMAHTNFHYHAE SQQQTTSTERRTTDSSTT HITRTVRORRAMTKELSSTT	
CKLLTGWCVTAKAPPDI LTAVKHGNCSSLTPLLNI VIVCAEMBEQWGYVGA RWLFYAYDSLRKTVVAI RTMATLGRLMSLSSPED MTDGWPLYESRLKGKLI KARSKOQKDADTAPPIQ TPCTYHAARDTROASPR QHRQADDDKRQTRHAE RTHTGRKTHKETATQQ, QTRRTARTHRRTARHTA RTRVAARQTTEKDNEQSS TQKTPSATTAHTPRRTI VQARHTSARERNHTRQH RETTTNAHTMPHYHAE SQQQTTSTERPTIDSSTT. HTTRTVROERRMGHEKI STTTNRTTTTGAYQHTQI	
LTAVKHGNCSSLTPLINI VIVCAEMDEQWGYVOX RWLFYAYDSLRKTVVAI RTMATLGRLMSLLSPFD MTDGWPLYESRLKGKLI KARSKQQKDADI'APPIQ TPQTVHAARDTRQASPR QHRQADDDRQTRHAE RTHTGRRKTIKETATQQ, QTRRTARTHRRTARHTA RTRVAARQTTEKOMEQS TQKTPSATATAHTPRRTI VQARHTSARERNHTRQH RETITINAHTHPHYHAE SQQQTTSTERPTDSSTT HTPRTVRDERRMTQHKI STTTNRTTTTGAYQHTQ	
VIVCAEMDEQWGYVGA RWLFYAYDSLRKTVVAH RTMATLGRLMSLLSPFD MTDGWPLYESRLKGKLL KARSKQKDADTAPPIQ TPQTYHAARDTRQASPR QHRQADDDKRQTRHAE. RTHITGRRKTIKEFATQQ. QTRRTARTHRRTARHTA RTRVAARQTTEKDNEQS TQKTPSATATAHTPRRTI VQARHTSARERNHTRQL RETITINAHTHTPHYHAE SQQQTTSTERPTTDSSTT HTRTVRDERRMTQHKL STTTNRTTTTGAYQHTQI	
RWLFYAYOSLRKTVVAH RTMATLGRLMSLLSPFD MTDGWPLYESRLKGKLI KARSKQQKDADTAPPIQ TPOTYHAARDTRQASPR QHRQADDDKRQTRHAE RTHTGRRKTIKETATQQ, QTRRTARTHRRTARHTA RTRVAARQTTEKDNEQS TQKTFSATTATHTPRTT VQARHTSARERNHTRQH RETTTMAHTNIPHYHAE SQQQTTSTERPTIDSSTT HTPRTVROERRMOHEKI STTTNRTTTTGAYQHTQ	
RTMATLGRLMSLLSPED MTDGWPLYESRLKGKLI KARSKQOKDADI'APPIQ TPOTVHAARDTRQASPR QHRQADGDKRQTRHAE. RTHTGRRKTIKETATQQ. QTRRTARTHRRTARHTA RTRVAARQTTEKOMEQS TQKTPSATATAHTPRRTI VQARHTSARERNHTRQH. RETITINAHTHPHYHAE SQQQTTSTERPTTDSSTT HTPRTVRDERRMTQHK.	
MTDGWPLYERLKGKLI KARSKQKDADTAPPIQ TPQTYHAARDTRQASPR QHRQADDDKRQTRHAE RTHITGRRKTIKEFATQQ, QTRRTARTHRRTARHTA RTRVAARQTTEKDNEQS TQKTPSATTAHTPRRTI VQARHTSARERNHTRQH RETITINAHTNPHYHAE SQQQTTSTERPTIDSSTT HTRRTVROERRMGHEKH STTTNRTTTTGAYQHTQ	
KARSKQKDADTAPPIQ TPOTYHAARDTRQASPR QHRQADGDKRQTRHAE RTHTGRRKTIKETATQQ, QTRRTARTHRRTARHTA RTRVAARQTTEKDMEQS TQKTPSATATAHTPRRTIT VQARHTSARERNHTRQH RETTINAHTNTPHYHAE SQQQTTSTERPTTDSSTT HTTPTVRDERRMTQHKI STTTNRTTTTGAYQHTQ	
TPOTVHAARDTROASRE QHRQADGDKRQTRHAE. RTHTGRRKTIKETATQQ, QTRRTARTHRRTARHTA RTRVAARQTTEKDMEQS TQKTPSATATAHTPRRTT VQARHTSARERNHTRQH RETITNAHTNPHYHAE SQQQTTSTERRTTDSSTT HTTPRTVRDERRMTQHKH STTTNRTTTGAYQHTQ	
QHRQADDKRQTRIAE RTHTGRRKTIKETATQQ, QTRATARTHRATARHTA RTRVAARQTTEKDNEQS TQKTPSATATAHTPRRTI VQARHTSARERNHTRQH RETITIAMHTNJYHYHAE SQQQTTSTERPTIDSSTT HTTRTVRDERRMTQHK	
RTHTGRRKTIKETATQQ QTRKTARTHRRTARHTA RTRVAARQTTEKDMEQS TQKTPSATATAHTPRRTI VQARHTSARENHTTRQI RETITINAHTNTPHYHAE SQQQTTSTERPTTDSSTT HTPRTVRDERRMTQHKI STTTNRTTTTGAYQHTQI	
QTRRTARHTARTARHTA RTRVAARQTTEKDNEQS TQKTPSATATAHTPRRTI VQARHTSARERNHTRQH RETITINAHTNIPHYHAE SQQQTTSTERPTIDSSTT. HTPRTVRDERRMIQHAY	
RTRVAARQTTEKDNEQS TQKTPSATATAHTPRRTI VQARHTSARERNHTQH RETTINAHTNTPHYHAE SQQQTTSTERPTIDSSTT HTPRTVRDERRMTQHKI STTTNRTTTIGAYQHTQI	
TQKTPSATATAHTPRRTI VQARHTSARERNHTRQI RETITINAHTINPHYHAE SQQQTTSTERPTTDSSTT HTPRTVRDERRMTQHKI STTTNRTTTTGAYQHTQI	
VQARHTSARERNHTRQH RETITINAHTNIPHYHAE SQQQTTSTERPTIDSSTT HIPRTVRDERRMTQHKI STTINRTTITGAYQHTQI	
RETTINAHTNIPHYHAE SQQQITSTERPITIOSITI HIPRIVROERRMIQHKI SITTNRITITIGAYQHTQI	
HTPRTVRDERRMTQHKH STTTNRTTTTGAYQHTQI	•
HTPRTVRDERRMTQHKH STTTNRTTTTGAYQHTQI	INNNN
22504 52872 A 22632 103 313 ATAGCPSCSATEGVVRN	GKST
AGHORYLCSPCRKTWOL	
TASOP/GARIMGVGLNTV	
KNSGRSR	
22505 52873 A 22633 620 832	

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		Sequence		
22506	52874	Α	22634	453	1047	AADDRGHLRPTDGLLRNGKST
		l		1		AGHQRYLCSHCRKTWQLQFTY
		l				TASQPAGQ*RVTGTSERREQIIQ
		ĺ				RLRQQGSVQVNDLSALYGVST
ŀ		1				VTIRNDLAFLEKQGIAVRAYGG
		ı				ALICDSTTPSVEPSVEDKSALNT
		1				AMKRSVAKAAVELIQPGHRVIL
						DSGTTTFEIARLMRKHTDVIAM
		1				TNGMNVANALLEAEGVELLMT
						AGICALLTGCCVTAKAPPDISAI
						SALTAVKHGNCSSLTPLLNPPG
l .			1			SDVPVCAEMDEQWGYVGAKS
		1				RQRWLFYAYDSLRKTVVAHVF
		1				GERTMATLGRLMSLLSPFDVVI
1		l				WMTDGWPLYESRLKGKLHVIS
		l				KRYTQRIERHNLNLRQHLARLG
		l				RKSLSFSKSVELHDKVIGHYLN
						GDFLQSTLREAPLIPHTREHP
22507	52875	Α	22635	1089	1686	TSPGYWRANSTDPDGRYCRTG
		ĺ				TSGGVCHALSSLRCPSCSATDG
1						VVRNGKSTAGHQRYLCSHCRK
		ı				TWQLQFTYTASQPAGQ*RFLSK
		l				QEKTLLINQLASEEQQKVTSRL
						SDALRNGRVWQLAIIYLTIQVA
						VYGLIFFLPTQVAALLGTKVGF
						TASVVTAIPWVAALFGTWLIPR
						YSDKTGERRNVAALTLLAAGIG
						IGLSGLLSPVMAIVALCVAAIGF
Į						IAVQPVFWTMPTQLLSGTALAA
					1	GIGFVNLFGAVGGFIAPILRVKA ETLFASDAAGLLTLAAVAVIDR
		l				RGTSLNRLRNTMIALLSNPFKE
						RLRKGEVOIGLWLSSTTAYMA
		1				EIAATSGYDWLLIDGEHAPNTI
						QDLYHQLQAVAPYASQPVIRPV
		ı				EGSKPLIKQVLDIGAQTLLIPMV
		1				DTAEQARQVVSATRYPPYAVH
		l				PVQLLTGWCVTAKAPPDISAIS
						ALTAVKHGNCSSLTPLLNPPGS
		1				DVIVCAEMDEQWGYVGAKSR
1		l				QRWLFYAYDSLRKTVVAHVFG
1		1				ERTMATLGRLMSLLSPFDVVIW
		1				MTDGWPLYESRLKGKLHVISK
						RYTORIERHNLNLROHTGTGW
						DGSRCRSQKSGGAA
22508	52876	A	22636	91	200	SGIIAFQYLLMTLKVLPVASVSI
						SCPS\CSATDGVVRNGKSTAGH
		ĺ				QRYLCSHCRKTWQLQFTYTAS
		ĺ				QPAGQ*RGGA

22509 52877 A 22637 2649 3270 GAPVASVSISC NOKSTAGHQI LQFTYTASQP KNKHMTITEF GEISPINITAM LADIEAEDLA WRYLNRILLS VSYQTPTREIN RENGGIOQRW VPGSFNDQFA WYQRVFIPK DAVTHYGKV DAVTHYGKV DAVTHYGKV DAVTHYGKV	PSCSATDGVVR RYLCSHCRKTWQ AGQ*RTIGRAML VAQRMRDAQQK ITSRLLLDLTYGV RKRRLLRITRDD SLFRRAGDNCAR KKLDGLWAFSLD WESALQESRAIA DADIRNYAGNV
22509 52877 A 22637 2649 3270 GAPVASVSISC NOKSTAGHQI LQFTYTASQP KNKHMTITEF GEISPPINTAM LADIEAEDLA WRYLNRILLIS VSYQTPTERIN RENGGIOQRW VPGSFNDQFA WYQRVFIPK DAVTHYGKV DAVTHYGKV DAVTHYGKV DAVTHYGKV	PSCSATDGVVR RYLCSHCRKTWQ AGQ*RTIGRAML VAQRMRDAQQK ITSRLLLDLTYGV RKRRLLRITRDD SLFRRAGDNCAR KKLDGLWAFSLD KKLDGLWAFSLD DADIRNYAGNV
22509 52877 A 22637 2649 3270 GAPVASVSISG NGKSTAGHQI LQFTYTASQP KNKHMITTEF GEISPDINTAM LADIEAEDLA WEYLMRLILS VSYQTPTREIR RENGGIORW VPGSFNDQFA WYQREVFIFK DAVTHYGKV DAVTHYGKV DAVTHYGKV DAVTHYGKV DAVTHYGKV	RYLCSHCRKTWQ AGQ*RTIGRAML VAQRMRDAQQK ITSRLLLDLTYGV BKLFRLRITRDD BKLFRAGDNCAR KKLDGLWAFSLD /WESALQESRAIA DADIRNYAGNV
NGKSTAGHQI LQFTYTASQP KNKHMTITEF GEISPDINTAM LADIEAEDLA WRYLNRILILS 'VSYQTPTREIN RENGGIDQRW VPGSFNDQFA WYQREVFIPK DAVTHYGKV	RYLCSHCRKTWQ AGQ*RTIGRAML VAQRMRDAQQK ITSRLLLDLTYGV RKRRLRITRDD RKRRAGDNCAR KKLDGLWAFSLD /WESALQESRAIA DADIRNYAGNV
NGKSTAGHQI LQFTYTASQP KNKHMTITEF GEISPDINTAM LADIEAEDLA WRYLNRILILS 'VSYQTPTREIN RENGGIDQRW VPGSFNDQFA WYQREVFIPK DAVTHYGKV	RYLCSHCRKTWQ AGQ*RTIGRAML VAQRMRDAQQK ITSRLLLDLTYGV BKLFRLRITRDD BKLFRAGDNCAR KKLDGLWAFSLD /WESALQESRAIA DADIRNYAGNV
LQFTYTASQP KNKHMTITEF GEISPDINTAM LADIEAEDLA WRYLMRILLIS VSYQTPTREIF RENCGIDQRW VPGSFNDQFA WYQREVFIPK DAVTHYGKV DAVTHYGKV	AGQ*RTIGRAML VAQRMRDAQQK ITSRLLLDLTYGV RKRRLRITRDD SLFRRAGDNCAR KKLDGLWAFSLD WESALQESRAIA DADIRNYAGNV
KNKHMTITEF GEISPDINTAM LADIEAEDLA WRYLNRILLIS VSYGTYREH RENGGIOQRW VPGSFNDQFA WYQREVFIPK DAVTHYGKV	VAQRMRDAQQK ITSRLLLDLTYGV RKRRLLRITRDD SLFRRAGDNCAR KLDGLWAFSLD /WESALQESRAIA DADIRNYAGNV
GEISPDINTAM LADIEAEDLA WRYLMRILLIS VSYQTPTREIN RENGGIOREW VPGSFNDQFA WYQREVFIFK DAVTHYGKV	ITSRLLLDLTYĞV RKRRLLRITRDD SLFRRAGDNCAR KKLDGLWAFSLD WESALQESRAIA DADIRNYAGNV
LADIEAEDLA WRYNRILLIS VSYQTPTREII RENCGIDQRW VPGSFNDQFA WYQREVFIPK DAVTHYGKV	RKRRLLRITRDD SLFRRAGDNCAR KKLDGLWAFSI.D WESALQESRAIA DADIRNYAGNV
WRYLNRILLS . VSYQTPTREIN . RENGGIOQEW VPGSFNDQFA WYQREVFIPK DAVTHYGKV	SLFRRAGDNCAR KKLDGLWAFSLD WESALQESRAIA DADIRNYAGNV
. VSYQTPTREIK RENGGIDQRW VPGSTDQFA WYQREVFIRK DAVTHYGKV	KLDGLWAFSLD WESALQESRAIA DADIRNYAGNV
RENCGIDQRW VPGSFNDQFA WYQREVFIPK DAVTHYGKV	/WESALQESRAIA DADIRNYAGNV
VPGSFNDQFA WYQREVFIFK DAVTHYGKV	DADIRNYAGNV
WYQREVFIPK DAVTHYGKV	
DAVTHYGKV	
22510 52878 A 22638 II 2470 MNADTDYSIA	EAAFNKGETAM
	IDTSKVNYGVT
	KPFVGVLSAGIN
	EFLENYLLTDEG
	GAVALKSYEEE
	MENAQKGEIMP
	AVRTAVINAAS
	DAQTRITNCCRI
	RKNPMDVIKKKH
	SVLGLLGLLVG
	GEYLFAITTLILS
	KAYAWRYVYPG
	FPLVCTIAIGFNN
	MPAKSSSLRSSQ
	AERTRYDSPGSY
	RGGSPFGGPGKY
	IAPSAQNQSPHG
	HALDWLDAGYT
	PYFKLAYVYDDS
	SIDNGTEGSAVR
	KNFSAYTDANYL
	SANDLTGITAKD
	QEFGKLDKCLS
	EKQVIFSSDVNN
	SLRREENVLIGIT
	GLMKNDIPYKLI
	PAKQTITLIEQN
	LAWHNRILELRD
	QFAHIIFMIAAM
	FSSGQHSVARKS
	CHEGFTISANPNS
SPRYPTIIEQAI	EQPRIVDFSIVRF
22511 52879 A 22639 I 852	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
	1			sequence		
22512	52880	A	22640	Ī	747	MILIDPRYTDTGAGREDEWIPIR
		1				PGTDAALVNGLAYVMITENLV
						DOAFLDKYCVGYDEKTLPASA
		1				PKNGHYKAYILATD/GVVRTAK
						STAGHORYLCSHCRKTWOLOF
						TYTASQPAGQ*RYTDTGAGRE
						DEWIPIRPGTDAALVNGLAYV
						MITENLVDQAFLDKYCVGYDE
	1					KTLPASAPKNGHYKAYILATDG
	1				1	WCVPPKAPPDISAISALTAVKH
						GNCSSLTPLLNPPGSDVIVCAE
1		1				MDEOWGYVGAKSRORWLFYA
						YDSLRKTVVAHVFGERTMATL
						GRLMSLLSPFDVVIWMTDGWP
						LYESRLKGKLHVISKRYTQRIE
						RHNLNLRQHLARLGRKSLSFSK
						SVELHDKVIGHYLNIKHYO
22513	52881	A	22641	1722	2463	NPRLAGSEGSCPSCSATDGVVR
						NGKSTAGHQRYLCSHCRKTWQ
						LQFTYTASQPAGQ*RKLTPGCL
		1				FVALKGERFDAHDFADQAKAG
		1				GAGALLVSRPLDIDLPQLIVKD
			1			TRLAFGELAAWVROOVPARVV
		1				ALTGSSGKTSVKEMTAAILSQC
				1	l l	GNTLYTAGNLNNDIGVPMTLL
		l				RLTPEYDYAVIELGANHQGEIA
						WTVTHPPRFVTKQWLCFLPDG
		ı				LLLFAEARFVIWQMQPAVMAV
		1			i	DVAAFAHLFNQQQYQRFTGRP
		1				PDSSKLPWKITCNGECVTFHW
	1	1			i	WLLGVAVRISRLLPVRVMTAL
1		1				EGNYNGTEGISALPFNGIILAHS
l	-	1			l	NESEWVTFRNNKNNEAFLDRV
	1	1				YIVKVPYCLRISEEIKIYEKLLN
		1				HIFECWQHFLNNPESLQCYIWA
ļ		1			l	ANADFREARQVIFKTFFTVGVE
ł		ı				EELGIRQTRTNDFLVTGDNLLRI
		1				FRFDVGNEDKVRQQFAVVRIH
	ŀ	1				REVFLVTFHGVNQRFSRHREEF
1		1				LFEFCVSVTANLTRCFFGIGFNL
	ŀ	1				RLTFFVIGDDFPTLTQNFRILIGV
1	1				i	IDGEFRLAHKAVAANHAIGLNA
1						QNGCRNDFVAHAAGSRALAAD
	1	1			1	GRSLEHAPTVETVVAGPLCESG
						DVFTQQEGGNVETRALPEVKA
					1	AVPPVQLLTGWCVTAKALPDIS
						AISALTAVKHGNCSSLTPLLNPP
1	1				1	GSDVIVCAEMDEQWGYVGAKS
	1				I	RQRWLFYAYDRLRKTVVAHRL
			-			

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22514	52882	Α	22642	90	1254	KGLCSHSEMVVPSYSATDGVV
		l				RNGKSTAGHQRYLCSHCRKTW
	İ	1				QLQFTYTASQPAGQ*RDIVLTK
		ı				LMMVWHSQSGYEKAYAHIQR
		1				WWYRAILLLTGWCVTAKALPD
		l				ISAISALTAVKHGNCSSLTPLLN
	ì	1				PPGSDVIVCAEMDEQWGYVGA
		l				KSRQRWLFYAYDRLRKTVVAH
	1	l				VFGERTMATLGRLMSLLSPFDV
	1	1				VIWMTDGWPLYESRLKGKLHV
						ISKRYTQRIERHNLNLSSTSILVL
		1				LNGATQFQDLGDLNKELEETY
		1				TDSKNHISCVGNDFGFNDIFSR
						YVEAVGREGDVLLGISTSGNSA
						NVIKAIAAAREKGMKVITLTGK
		1				DGGKMAGTADIEIRVPHFGYA
		l				DRIQEIHIKVIHILIQLIEKEMVN
		l				FTGLVQRGGGTGPHKDGWGIT
						FYEVAFEDVAVNFTQEEWALL
		l				DPWQKKLYRDVMLETYRNLAS
		l				VGDDDNIPSLREQVAHQRYFKT
		<u> </u>			1000	WHVEREYFSK
22515	52883	Α	22643	690	I233	RLQSLHYQGRGKPVPMRSYSLS FLFTISNSAGVFRCPSCSATDGV
	1					VRNGKSTAGHORYLCSHCRKT
	İ	l				WOLQFTYTASQPAGQ*RLLGP
		l				KNRPREQNWLLWCERGVMSRL
		l				OALYLREOGENNVKVDNFOTK
						VDSCTRPLAGQQFYPCIPDDAL
		l				VOPLEGGNEVFORREAGHITHE
			ļ			EDIVRHQHGGWRGANSRDDTP
1				l		GTVELFNGFNOOSAIAKTLRAF
		l				NATROHDHVIFAISHFDORRIR
İ						QQLHAARAGNRQVTIAGDAGS
		l				GDFDTAANQQIDSGDGFSLFTT
	1	1				RGEANQCRCAHIHSPFFLPFQT
			1			VRVFSAVPPVQLLTGWCVTAK
		1				APPDISAISALTAVKHGNCSSLT
		1		1		PLLNPPGSDVIVCAEMDEQWG
	1	1		1		YVGAKSRQRWLFYAYDSLRKT
						VVAHVFAVKPYWDTGGGFHC
	Ī	-		1		RWIYMQQEKELLQLSLQQGKY
	1		t			NOKRAAELLGLTYHOFRALLK

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22516	52884	A	22644	1314	1483	RIQRYCGKTGCPSCSATDGVVR KOKSTGAHORYLCSHCRKTWQ LQFTYTASQPAGQ*RMFNADG SEVAQCGNGARCFARFVLKG LUTNKRDIRVSTANGKMVLTVT DDDLVRVNNGEPNEFPSAVPF RANKAEKTYMRAAEQFILCGV VSMGNPHCVIQVDDVDTAAVE TLGPVLESHERPFERANIGFMQ VVKREHIRLRVYERGAGETQA COSGGACAAVAVGIQQGLLAEE KYVELPGGRIDJAWKGEELQET LTELDDRAVVDYLIKNPEFFIRN RAAVEAIRVPHPVGGTVSLVE WHMARARNHIHVLEENNALL MEQAJANEGLFYBLLYLQRSLT LAYAMPVTINKGNQIENQLDT ETITISYEFILCOHLIANTEFSYL ALPENYNRLCHPNSKNQTNNRF KTLNSKAIGRILAAGGVYNGNI EGRFTATEKLAVPPVQLLTGW CVTAKALPPISAISALTAVKHG NCSSLYTASOPAGO

NO:	of peptide sequence		SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	deletion, \=possible nucleotide insertion)
222517	52885	A	22645	1479	1793	TRYRSCPSCSATDGVVRNGKST AGHQRYLCSHCRKTWQLQFTY TASQPAGQ*RRNACQYRTPAV VASRQKRWHGOLGMIGSROV MLFISDSVGAKELDLIPRLEDK SIALLAMTGKFTSPLGLAAKAV LDISVEREACHMILAPTSSTVN TLMMGDALAMAVMQARGFNE EDFARSHPAGALGARLLNKVH HLMRRDDALPOVALTASVMDA MLELSRTGLGLVAVCDAQQAV QGVFTDGBLRRWLVGGGARLLNKVH PVNEAMTVGGTTLQSQSRAIDA KEILMKRKITAAPVVDENGKLT PVNEAMTVGGTTLQSQSRAIDA KEILMKRKITAAPVVDENGKLT VSHDRHRDAERQRVKSGLPRY AGAGLANRARNWYGSSWK VOYGOJILQGGPQNAEKSGQRVT VSHDRHRDAERQRVKSGLPRY AECSTYPSCPVALDLAFASHGI HMHAPGEALRMLVTAMDKAA DARTKLARLLATKGITHEIQIPD ISTKEKAQQFIGLNMEQIKPEKQ DFIKPVIPQAGKHNGVNMSVLR SLLTAGVLASGLLWSLNGITAT PAQASDDRYEVTQQRNPMSVLR SLLTAGVLASGLLWSLNGITAT PAQASDDRYEVTQQRNPMSVLR SLLTAGVLASGLLWSLNGITAT COYTAKAPPDISAISALTAVKHG CVTAKAPPDISAISALTAVKHG CVTAKAPPDISAISALTAVKHG CVSSLTPLINPPGSDVIVCAELD EQWGVYGAKSRQRWLFYAYA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22518	52886	Α	22646	807	1350	NRGCPSCSATDGVVRNGKSTA
				1		GHQRYLCSHCRKTWQLQFTYT
						ASQPAGQ*RLLNRLIEPTRGQV
						LIDGVDIAKISDAELREVRRKKI
						AMVFQSFALMPHMTVLDNTAF
						GMELAGINAEERREKALDALR
				l		QVGLENYAHSYPDELSGGMRQ
						RVGLARAINPDILLMDEAFSAL
		1				DPLIRTEMQDELVKLQAKHQR
						TIVFISHDLDEAMRIGDRIAIMQ
				ŀ		NGEVVQVGTPDEILNNPANDY
					l	VRTFFRGVDISQVFSAKDIARRT
						PNGLIRKTPGFGPRSALKLLQD
						EDREYGYVIERGNKFVGAVSID
						SLKTAAVPPVQLLTGWCVTAK
						APPDISAISALTAVKHGNCSSLT
						PLLNPPGSDVIVCAEMDEQWG
						YVGAKSRQRWLFYAYDSLRKT
						VVAHVFGERTMATLGRLMSLL
				i		SPFDVVIWMTDGWPLYESRLK
		l	ĺ			GKLHVISKRYTQRIERHNLNLR
						QHLARLGRKSLSFSKSVELHDK
22519	52887	A	22647		1270	VIGHYLNIKHYQ MTAATEOLNASRGLSTREMER
22319	32007	^	22047	l'	1270	OAELAKITTDYINSGGSEGDEK
						LONMIKAONDYYAAEDAKRA
						DWLAGAESAFADYGDSAMDM
						YGNVNEIASSALNGMTDMMV
						OFLTTGKANFADFAKNIISMIIK
						MIAOMVIFNTSLRNDGRSVSSG
						GAAVSLAAAAGSVATSGFNAS
		l	1			NSAPKVVTIREVERSVDVSGME
						ASSTSWITIRSFTAETSTFGPTM
						TASMTFLPTYCSIIASLLSSFKLA
						AATDGVVRNGKSTAGHQRYLC
i		i	}			SHCRKTWQLQFTYTASQPALA
		1		l		RNPNRDITIYWGGREEQHLYDL
		1	1			CELEALSLKHPGLQVVPVVEQP
						EAGWRGRTGTVLTAVLQDHAT
		1		l		DGVVRNGKSTAGHQRYLCSHC
						RKTWQLQFTYTASQPGTHQKII
				ĺ		DMAMNGVGCRATAR\IMGVGL
	l	l				NTIFRHLKNSGRSR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22520	52888	IA	22648	381	945	GAPVASVSISCPSCSATDGVVR
22320	32000	^	22048	381	243	
						NGKSTAGHQRYLCSHCRKTWQ
				i		LQFTYTASQPAGQ*RTENQTPH
		ı				VLTHKWELNYENPWTQGGEHL
		l				TMGLVVGWRGLQEGDPARTN
		l				LSYYQANVGSGQHRENKGPIN
		1				LGNVGKSNGERGGKSLYRCKIF
			İ	l		LESSVTMKDRELPPVYDGVFEV
		1	l			LQWLLFLSAVPPVQLLTGWCV
		l		l		TAKAPPDISAISALTAVKHGNC
						SSLTPLLNPPGSDVIVCAEMDE
		1	1	l		QWGYVGAKSRQRWLFYAYDS
						LRKTVVAHVFGERTMATLGRL
		1				MSLLSPFDVVIWMTDGWPLYE
				l		SRLKGKLHVISKRYTORIERHN
						LNLROHLARLGRKSLSFSKSVE
		ŀ		1		LHDKVIGHYLNIKHYQ
22521	52889	A	22649	354	1119	RRWRRGAHEAATGGTGRGPAE
22321	32007	l^	22047	1234	1117	EAAPGEPAAGRAAPKSGGKRL
						GGVLGRGGVGRRSGSPDRSGA
						HSAAPRRCRQLPGRCPSCSATD
			l			GVVRNGKSTAGHORYLCSHCR
						KTWQLQFTYTASQPAGQ*RSK
						GAIGEGPPDPRGIVEQNLGGDIS
	1				ľ	IPCYNGPRPRRSAQKTTLGPLN
				l		QPRIPSAQCAGAFPRVAAAGAG
				l		GGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG
	1	1		l		RQRRRRWRRGEGEGDGDAGH
	1					MRPRPAGRAEARRRRRLRGNP
		1				PPGARLPNREGNGSGVSWAAE
						VSGDGRGARTEAARTRRPGG
				l		ADNYPGAVPPVQLLTGWCVTA
	1			1		KAPPDISAISALTAVKHGNCSSL
						TPLLNPPGSDVIVCAEMDEQW
					ľ	GYVGAKSRQRWLFYAYDSLRK
						TVVAHVFGERTMATLGRLMSL
				l		LSPFDVVIWMTDGWPLYESRL
						KGKLHVISKRYTQRIERHNLNL
						ROHLARLGRKSLSFSKSVELHD
		l				KVIGHYLNIKHYQ
22522	52890	A	22650	23	658	VISRPGSSIQVRGLLLSRHGLAL
		Ľ			1	GSAPDLWKLPWVNLVQGMGQ
		1	l			OSOGSSCREGCSRPOCPSAAPC
1	1	1	l	I		GAAPAHFCSAATFWESEVQSIP
	1	1	1	I		R/CRATDGVVRNGKS/TAGHQR
				1		YLCSHCRKTWQ/LQFTYTASQP
	1	1		I		
		1		I		EVIRSSPTSAPRSDGTKTVPPLM
		1	1	1		GISP*SAHHRLATSDPSLCLSES
	1	1	1	I		LCSREPA A VPRASNQQPQGVK
		_				DVHHQFTARTSRYPSKC
22523	52891	A_	22651	926	1093	

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22524	52892	A	22652	343	817	GGARSRQNPRCGPCMVVVPGQ
22324	32072	^	22032	545	017	SGRSVSADSSTEFREQGGS\AI\S
		l				FPKSINRAGPELFRFMENPFNSV
		l				*YRKNVRELTPAIMAVLPVEFR
				i		TRLAPONDTMSLIASAMKECSE
		ı				AKQAVLLNAPEHQKMKEVSEG
		l				IASLFRLMPEQVGPLMTMVTS
						MLGVI
22525	52893	A	22653	115	425	111111111111111111111111111111111111111
22526	52894	A	22654	747	1173	
22527	52895	Α	22655	I	2709	
22528	52896	A	22656	1	2583	
22529	52897	Α	22657	636	1029	APSDQKRDFLVPHGADSAMAK
						SG*CESGNDLLSSSHMNQLR/A/
		l				ESIPEDVIAGASALVLTFIIKFVI
						AENPQWWQQFLKDHVSILAMN
						EDEAEALTGESDPLLASDKALD
						WVDLVLCTAGPIGLYMAGGGD
						VN
22530	52898	Α	22658	1611	3300	QSAWANRHRLVDCGRPGNGSD
						AGVAARSGRNDGTGRCGLCHS
						CSRYGDHHRQSDRIYRHYDAG
		1				RSPSGAVDYGTQRGNRFSRAV
						YPVGA/RRWR*GLPLV/QQFQPE
		1				NETSAAW\VVGINQPLVDIEAK
						VDDEFIERYGLSAGHSLVIEDD
				ł		VAEALYQDVQQKNLITHQSAG
						GTIGNTMHNYSVLADDRSVLL
				١.		GVMCSNIEIGSYAYRYLCNTSS
				l		RTDLNYLQGVDGPIGRCFTLIG
	l					ESGERTFAISPGHMNQLRAESIP
		l	ĺ			EDVIAGASALVLTSYLVRCKPG
		İ				EPMPEATMKAIEYAKKYNVPV
		l			ŀ	VLTLGTKFVIAKNPQWWQQFL
					1	KDHVSILAMNEDEAEALTGESD
				1		PLLASDKALDWVDLVLCTAGPI
			l			GLYMAGFTEDEAKRKTQHPLL
		ĺ				PGAIAEFNQYEFSRAMRHKDC
						QNPLRVYSHIAPYMGGPEKCNI
			l	1		LGRSGWLLYGRGLDKQRLTQY
		ĺ		1		QSKLGAAMVIVAAWCVEDYQ
	1		l	1		VIRLAGSLTARATRLAHEAQLD
	1		1	1		VAPLGKIPHLRTPGLLVMDMDS
				1	1	TAIQIECIDEIAKLAGTGEMVAE
						VTRTGDARRTRFYRQPAQPCG
22531	52899	Α	22659	430	586	
22532	52900	Α	22660	1	131	

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1		l		sequence		
		_				
22533	52901	Α	22661	ı	I282	MLPAHLDACVTTLQKIRWTLS
					ŀ	VVDPTLALRVPLPALSIRDDPPL
						RAYLPGRGALKVIRFDDMDFA
		l			ŀ	TIDGINRMLVDIDTNYLFLPRVT
		l				SSTIRCLSVPTRRTVPAVTASGR
						SVKYYPPDFDPSKIPKLKLPKD
						RQYVVRLMAPFNMRCKTCGEY
					1	IYKGKKF\NARKETVQNEVYLG
	i	l			l	\LPI\FRFYIK\CTRCLGRDSPFKT
1					1	DP\ENTDYTM/ESLGATRNFOG*
						EASWEEEEKRV\QK\EREDE/ES
						LNNPMKVLENRTKDSKLEMEV
						LENLQELKDLNQRQAHVDFEG\
						MLRQHRLSEEERRSQQQEEFEQ
					l	KTAALLEEARKRRLLEDSDSED
ł						EAAPSPLOPALRPNPTATLDEA
						PKPKRKVEVWEQSVGSLGSRPP
1						LSRLVVVKKAKADPDCSNGQP
l						QAAPHPRSPAEQEGGQPYTPDA
						WRVLPEPTGCIPGQ
22534	52902	A	22662		412	WKVEFEFTGEFGQ
22535	52903	A	22663	298	532	MGSWSLPGRTLWKEKVRFLKS
22333	32303	n	22003	270	1552	KERTNPGSNFFSFPGSQHTREER
		1	l			SQECLSLPVLPSYLATRLSLPDS
		1	l		1	CPLKGRKAH*QRHSEI*LMGFE
		1			i	PGWPGKL*GWIELAFLLGCLHL
į.					1	TLPLPCPTPLPPKSTKSGCFSLSI
		ļ.				ALCFNKQFAVTK
22536	52904	A	22664	1	193	ALCFNKQFAVIK
22537	52904	В	22665	55	281	
22538	52905	A	22666	86	159	
22539	52907	A	22667	63	1036	VRGGLAAGRGRGSAGAAPVVV
22339	32907	A	22007	03	1030	AAMLGAWAVEGTAVALLRLL
		l				
i		l				LLLLPPAIRGPGLGVAGVAGAA
		1				GAGLPESVIWAVNAGGEAHVD
		1				VHGIHFRKCIPL*GRVGRALYY
		1				GMKLPILRSNPEDQSCYQTE\RS
		l				N/EETFGYESPIQREGGLTCWSL
		l				KFAE\VYFAQSQQKVFDVRLNG
		l	l	1		HVVVKDLDIFDRVGHSTAHDEI
		1	l		ĺ	IPMSIRKGKLSVQGEV\ST\FTQG
		1				KLLPLSFSKGYL*PIPKVCAL\YI
1	1	1	l	l	l	MAG\TVDDVPKASAFNPGLEK
		1	l	ĺ	i	KEEEEEEEYDEGSNLKKQTNK
		1				NRVQS\GPRTPNPYALDNSSLM
L		L_				FPILV\AFGVFIPTLFCLCRL
22540	52908	Α	22668	1570	1773	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
	L	丄				L
22541	52909	A	22669	62	454	NKQLNKKRRRTMIALIQRVTRA
		1				S/VTVEGEVTGEIGAGLLV/LLG
						VEKDDDEQKAN/RDAEGQM/N
		1				LNVQQAGGSVLVVSQF/TLAAD
		1				TERGMRPSFSQGHHRIRAQAFY
	1	1		i .		DYFVES/CRQQR*YAQGGLSAE
						FRLLNS
22542	52910	A	22670	1	1930	MAGKPEYDKTISTSIVLNALNA
	122710	ľ.		ľ		LGVSAEASGRNDLVVKTVEGD
		1				RKVSGSAYRETKDRGFHHGTL
		1	Į.	İ		LLNADLSRLANYLNPDKKKLA
				ł		AKGITSVRSRVTNLTELLPGITH
	1	1	ì	j		
İ			l	ĺ		EQVCEAITEAFFAHYGERVEAE
	l	1	ĺ			IISPNKTPDLPNFAETFARQSSW
						EWNFGQAPAFSHLLDERFTWG
		1		ŀ		GVELHFDVEKGHITRAQVFTDS
1		1				LNPAPLEALAGRLQGCLFSLYN
		1				SDLNTVIDNVLRIFPLLLSWISF
		1				WLLYSIVPTIRVPNRDAIVGAFV
	l	1			}	AALLFEAGKKGFALYITMFPSY
						QLIYGVLAVIPILFVWVYWTCV
	l	1	l			TVEGEVTGEIGAGLLVLLGVEK
				1		DDDEOKANRLCERVLGYRIFSD
l	l	1	1			AEG\KMNLNVQQAGGSVLVVS
İ	l	1				OFTLAADTERGMRPSFSKGVID
	1	1				DIGFNTEISQTRFHQASHLFKRK
	l	1				GIYRLHFARRRCQQIQRRQRRR
		1		l		RFMORTAVAITHOORMKOGIL
		1				RLWRNWLLWLLILLLRRNRLL
	l	1		ŀ		RLLILLLWRNRLLRLLILWLRR
	1	1		1		NRLLWLLILRLRRNWLLRLLIL
						LLRCNRLLRLLYRFNNWRKQR
						FVWTIIQQRFHWRKFKIIQRERI
						RPSRLFAGITLLLSLRKLTSQFS
		1	l			LCRSICIFCVYGHIMLILFAIALL
						LCLGELTCQFILLHRIDFIIVVLI
22543	52911	A	22671	377	463	QTISQTTVQSN*NSGLRISPKTA
		1				QLHGD
22544	52912	Α	22672	129	615	KARQKSCQPVTTDTQCRHTSSS
1		1	1			IGKRLAFCQNLTQIHTESTMKP
1	1	1	ĺ	1		WILLLVMFISGVVMLLPVLGSF
1	1	1		1		WNKDPFLDMIRETEQCWVQPP
		1		1		YKYCEKRCTKIMTCVRPNHTC
1		1		1	1	CWTYCGNICLD/IRRAP*/HQC*
l						TPRFYWPITTVG*RMTIWSKKS
1		1		1		
22515	52012	١.	22672		412	AAYFLLGTPTWL
22545	52913	Α	22673	3	412	
22546	52914	Α	22674	2	390	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop eodon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	1			sequence		
22547	52915	Α	22675	180	507	CCGGRARSELCLLNLCIKGAD*I
		l		1		QAW*KFCQWPCERADGRGCRK
						AER\CPCPKGAYSL/YCRESVCC
						LPSFVNKNSSSLCVAHNWGRH
1		l				CYLWTAADVKQFISSHMFPVE
1						HL
22548	52916	Α	22676	3	227	GGGRARSELCLLNLCIKGAD*I
		1				QAW*KFCQWPCERADGRGS*F
		ı				DRGAEVCNQTDWQSIRSSQGG
						RDQRFSTERA
22549	52917	A	22677	567	842	
22550	52918	Α	22678	1	1400	MGWGQGPDVQLMNGHERSPS
		1				HSPREAKKSGSPLPTDIHCFSPQ
1		ı				SQPAMVITGVPTAPAGASAKM
1						GLCATPITGACPLWLRGFRGWR
ł		1			İ	CEDRCEQGTYGNDCHQRCQCQ
1		1		1		NGATCDHVTGECRCPPGYTGA
						FCEDLCPPGKHGPQCEQRCPCQ
			ŀ			NGGVCHHVTGECSCPSGWMLS
					1	FPGWRPI*FSKSL*MQGTVCGQ
	1			l		PCPEGRFGKNCSQECQCHNGGT
		l		l		CDAATGQCHCSPGYTGERAAV
						PDVRK\CQDECPVGTYGVLCAE
l		l			1	TCQCVNGGKCYHVSGACLCEA
	İ	1		l		GFAGERCEARLCPEGLYGIKCD
						KRCPCHLENTHRGDDREKAEF
				i		GHLLIALEVQYTEISNCIEKSQE
		1				ASCYDIRMLKQPYGETNTEEEL
1	1			1		AVNYVYSIYASKEQIRYKHGRS
1	1					FANGHVKELMEGPSRRIAGSGG
l						EKYKRDEEKNTLLRGQNRGGR
1				1		GKNPKKTKAWVKIQGQNKIEIS
		L				GVGADRKRFFREK

SEQ ID SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	location of first	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
52919	A	22679		2210	MEÖKGEFKNDTLIYEFFHIVAK PHPLPIKSSTYTTFQSARVTRFF LDAEKECKCQEGQVEAALIAM GGTCQGTQRLSTAGESESATH MREPAVQCRENSOCYSVLAGR RRFFQGAASPLHPASVALTQLH VRWASTATIGQVSGFPGKIRVS PTINLGDSDYIFTASHWGLGHY GPAVHVHSQGGIDERGSNDSG RLYVTNPGGNPSWTPNIDRLA EDGVKLTQHISAASLCTPSRAA FLTGRYPVRSGMYSSIGYRVLQ WTGASGGIPTSETTFAKILKEK GYATGLIGKWHLGILNCESASD HCHHPHLHGFDHFYGMFPSLIK GDCARWELSEKRVNLEQKLNF LFQVLALVALTLVAGKLTHLISEY VGALIVHADCELMRNHTITEOP MCFQRTTPLIFLQEVASFLKRNK HGPFLLFVSLHVHIPLITMENF LGKILDFILDYGDNVSEMLDWFNS GGKSLMOLGNTQGWAGIY KGGKOMGGWEGGIRVPGIFRW PGVLPAGRVIGEPTSLMDVFPT VVKLAGGEVPODRVIGQULT VITAGREYGDFTSLMDVFPT VVKLAGGEVPODRVIGQULT VITAGREYGEPTSLMDVFPT VVKLAGGEVPODRVIGGWKVCFC VLHAARWHQRDRGTMWKVHF VTPVPQPEGAGACYGKKVCPCF EKCVHIPPLLEDLSROPSET

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22552	52920	Α	22680	185	2091	GSTERDNMLHLHHSCVCFRSW
						LPAMLAVLLSLAPSASSDISASR
						PNILLLMA\DDHGVGDIGCYGN
	ŀ				i	NTMRTPNIDRLAEDGVKLTQHI
i	İ	l		İ		SAASLCTPSRAAFLTGRYPVRS
l						GMVSSIGYRVLQWTGASAGFT
					l	TN\ETTFAKILKEKGYATGLIGK
			İ			WHLGLNCESASDHCHHPLHHG
						FEHFYGMPFSLMGDCARWELS
						EKRVNLEQKLNFLFQVLALVA
						LTLVAGKLTHLIPVSWMPVIWS
						ALSAVLLLASSYFVGALIVHAD
		i				CFLMRNHTITEQPMCFQRTTPLI
						LQEVASFLKRNKHGPFLLFVSF
1		İ		ŀ		LHVHIPLITMENFLGKSLHGLY
		i				GDNVEEMDWMVGRILDTLDV
						EGLSNSTLIYFTSDHGGSLENQL
		1				GNTQYGGWNGIYKGGKGMGG
1		1				WEGGIRVPGIFRWPGVLPAGRV
1						IGEPTSLMDVFPTVVRLAGGEV
ļ						PQDRVIDGQDLLPLLLGTAQHS
ŀ						DHEFLMHYCERFL\HAARWHQ
l						RDRGTMWESPTL*PPCVSSQEG
						AGAC\YGRK\VCPCFGEK\VVHH
			ŀ			DP\PLL\FDLSRDPSETHILTPASE
		l			1	PVFY\QVMER\VQQAVWEHQR
		1				TLSPVPMQLDRLGNIWRPWMQ
ì		l				PCCGPFPLCWCLREDDPQINVC
		L				SEKLEPRFLNFVTQIETNQLAM
22553	52921	Α	22681	209	392	
22554	52922	Α	22682	1	513	
22555	52923	Α	22683	3	613	
22556	52924	A	22684	1	2115	

NO:	SEQ ID NO: of peptide sequence	Mei hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22557	52925	A	22685	53	1966	VGQPRGLRTLEKLTASVAGGA ANFFGDIERDLGPRDTSRGIRG GNLGDRGSSAEMAYASWQR WSPVEWARWAWHATAYTSSGDS SLLVLQGDSGKRSSDSEEAFET EYSTTPVKASPAPPPPPPEVIPE EVSTGPPEEPGWGSETVPVPD GPRSDSVEGSPFRTOAHSFSGVF DEDQPIASTGTDILDFDNIELVD TFQTLEPRASDAKNQGGKVNT FRKSTIGSVPISKSTLSRSLSLQA SOFDGASSSGNPEAVALAPDAY STESAKRVVPPASGGGRVQNSP PVGRKTLPLTTAPEAGEVTPSD SGGQEDSPAKGLSVRLEFDVSE SUSSWDNQGENPPTKKIGKKP VAKMPLRFRMKKTPEKLDNT PASPPRSPAERDIPHKGTYTIF DIDKWDDPNFNPPSSTSKMQES PKLFQQSYNTDPDTCDESVDPF KTSSKTTSSFSKSPASFEPBASA MEANAVDGGGLNKPAKKKT PLKTMVEDVMSVCSLEDTFRY KKSPKRSPLSEDPSQDFTPAATP ETPPVISAVVHATDEEKLAVTN GKWTCMTVDLEADKQDYPPAATP ETPPVISAVVHATDEEKLAVTN QKWTCMTVDLEADKQDYPQPG
						YEIEYMEKIGSSLPQDDDAPKK QALYLMFDTSQESPVKSSTVR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
22558	52926	A	22686	662	2927	HLRRKRNLPRREGLSPALLEET
						PLEPAVGPKAACPLDSESAEGV
l	i					VPPASGGKRCLFTTAPEAGEVT
		1				PSDSGGQEDSPAKGLSVRLEFD
	ŀ	ŀ				YSEDKSSWDNQQENPPPTKKIG
						KKPVAKMPLRRPKMKKTPEKL
		l				DNTPASPPRSPAEPNDIPIAKGT
	l		İ			YTFDIDKWDDPNFNPFSSTSKM
		ŀ				OESPKLPOOSYNFDPDTCDESV
1		1				DPFKTSSKTPSSPSKSPASFEIPA
1	i	1				SAMEANGVDGDGLNKPAKKK
		l				KTPLKTDTFRVKKSPKRSPLSD
	i	l				PPSQDPTPAATPETPPVISAVVH
	ŀ	l			1	ATDEEKLAVTNOKWTCMTVD
		l			İ	LEADKODYPOPSDLSTFVNETK
	İ	l				FSSPTEELDYRNSYEIEYMEKIG
		l				SSLPODDDAPKKOALYLMFDT
i .	İ	l				SQESPVKSSPVRMSESPTPCSGS
		1	i			SFEETGSPLRNTAA\KNO\HPVP
		l			İ	TRDWAP*PRSSHLAGCQRKSSQ
		1				E/VELGGPWGLGHPFQEGDLKF
1		1				TASPRGSFASADALLSRLAHPV
		1				FSL/CGALDYL\EPNFRQKRNPP
		Ì				LFAQKLQREA\VHPTDVSI\SKT
1			l			ALYSRIRTTEVEKPAGLLFQQP
ł					i	DL\DSALQIARAEII\TKEREVSE
		l			İ	WKDKYEESR\REVMEMRKI\LA
					i	EYEKTIAQMIEDEQREKSVS\HQ
						TVQQLVLEKEQA\LADLNSVEK
				1		SLADLFRRYEKMKEVLESFRK
		l		1		NEEVLKRCAQEYLSRVKKEEQ
						RYQA\LKVHA\EEKLDRANAE\I
22559	52927	Α	22687	374	1268	AETNMVWYDWMRPSHAQLHS
						DYMQPLTEAKAKSKNKVRGV
		1		1		QQLIQRLRLIKSPAEIERMQIAG
				•		KLTSQAFIETMFTSKAPVEEAFL
		i				YAKFEFECRARGADILAYPPVV
		l				AGGNRSNTLHYVKNNQLIKDG
		i				EMVLLDGGCESSCYVSDITRTW
						PVNGRFTAPQAELYEAVLEIQR
	l		1		l	DCLALCFPGTSLENIYSMMLTLI
						GQKLKDLGIMKNTQGK*CL/RR
			İ			LLE/QYCPHHVGHYLGMDVHD
1		1	İ			TPDMPRSLPLQPGMVITIEPGIYI
			1			PEDDKDAPEKFRGLGVRIEDDV
	1	1				VVTQDSPLILSADCP
					L	

SEO ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22560	52928	IA	22688	329	1389	RIWKTOMMRMKKMMMMMM
22560	52928	A	22088	329	1389	
				i	1	MMMRTTLFFQQESQEAIFFPFD
		İ		1		LFPMCPFGCQCYSRVVHCSDLG
						LTSVPTNIPFDTRMLDLQNNKI
		1				KEIKENDFKGLTSLYGLILNNN
1						KLTKIHPKAFLTTKKLRRLYLS
						HNQLSEIPLNLPKSLAELRIHEN
					}	KVKKIQKDTFKGMNALHVLEM
i						SANPL**/SMGIEPGAF*RG*RCS
						/YIRIAEAKLTSV\PKGL\PPTLLE
						LHLDYNKISTVELEDFKRYKEL
					'	QRLGLGNNKITDIENGSLANIPR
				ŀ		VR\EIHLGNIKLKKIPSGLPELKY
						LQIIFLHSNSIARVGVNDFCPTV
i						PKMKKSLYSAISLFNNPVKYWE
						MQPATFRCVLSRMSVQLGNFG
22561	52929	Α	22689	1	4608	
22562	52930	Α	22690	1	465	
22563	52931	Α	22691	1	1542	
22564	52932	Α	22692	1	4226	EAVLCSSRGRPDSSMPDCCRNA
		1				ALEARSCQSMTSLFSNTVSPTQ
		1			ŀ	DGTSSLPRRQSSFAKPPLRALY
		1	l			DLLIAPMEGGVVESTEPLGTSR
		ł				SGFASGTALFWLGDNGKAASP
1		l				PGVSVSSLENEDDNTQCNRLEQ
						GYREAAVPDWTODLALCLLPT
		l				LOSLKARTAPHLCVPSSCYKA
		1	1			WYTLAVSKVSGMVEGKAWIIL
		1				LLLAASTHALLQRPQEGDITPL
		ı				AGSRAAAVSCOARPFSTFSGLM
		ı				HSSGPVGRHROLILVLE
22565	52933	A	22693	2	4197	ALDFPGRRFRPTASFIWASVLFE
22303	32,755	ľ`	22073	ľ	4157	TIRHEAEVSTDYKLSLFDLQTSS
		l				YQALQRVLVSLGHHDEALAVA
	1	ı	1			ERGRTRAFADLLVEROTGOOD
		ı				SDPYSPVTIDOILEMVNGORGL
		ı				VLYYSLAAGYLYSWLLAPGAG
		ı				
		l	1		1	IVKFHEHYLGENTVENSSDFQA
		l		1		SSSVTLPTATGSALEQHIASVRE
		I	1	l		ALGVESHYSRACASSETESEAG
		ı				DIMDQQFEEMNNKLNSVTDPT
		1		l		GFLRMVRRNNLFNRSCQSMTS
		L				LFSNTVSPTQDGTSS
22566	52934	Α	22694	1	618	
22567	52935	Α	22695	1	3528	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
22568	52936	A	22696	85	1193	RPIEGOLGTCHPGSEPRTIRWEA PGKGRLRVTHPPTSDIAGRTGA AASFPAPLHLPREPRAGVDGTD AERILYPRRAAPAASGVOSRPA GEAVRADEAGAAGLGGFQSER NRSWGGESLGSRQPOKMPAFN RLPPLASLVLIV*VSVCEPPWG VPLENGGRARANPMKLRCISC MKREEVEATTVVEWFYRPEGG LQWNGSKDLQDVSITVLNVTL NDSGLYTCNVSREFEFEAHRP VKTTRLIPLRVTEEAGEDFTSV VSEIMMYILLVELTLWLLEMIY CYRQVSKAEEAAQENAFGNSL EGGSQGLQNNSTQDGLVTGQQ DVGSHRAREHMPDFEEPPSAK
22569	52937	A	22697	277	527	VRVSWEYFCAPGPAGSPAGPG RGEEKQRLGLLIRDARSPAPPG PPVGSVRG/PKPAPAGRVRAKG RRA*GAACSLAAGAWVEPGI

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	· ·			sequence		· · · · · · · · · · · · · · · · · · ·
22570	52938	A	22698	226	2268	SVKNYTKCHVRNEQICNKLTSC
	1		l		1	KSCSLNLNCQWDQRQQECQAL
		ı	ŀ			PAHLCGEGWSHIGDACLRVNSS
		1	ĺ		1	RENYDNAKLYCYNLSGNLASL
		l			l	TTSKEVEFVLDEIQKYTQQKVS
						PWVGLRKINISYWGWEDMSPF
		l				TNTTLOWLPGEPNDSGFCAYLE
				ŀ		RAAVAGLKANPCTSMANGLVC
		l				EKPVVSPNONARPCKKPCSLRT
		l	ļ			SCSNCTSNGMECMWCSSTKRC
		l	ŀ			VDSNAYIISFPYGQCLEWQTAT
		l		l		CSPONCSGLRTCGOCLEOPGCG
		l	ŀ			WCNDPSNTGRGHCIEGSSRGPM
						KLIGMHHNEMVLDTNLCPKEK
					Į.	NYEWSFIOCPACOCNGHSTCIN
		1		İ		NNVCEOCKNLTTGKOCODCMP
		l				GYYGDPTNGGOCTACTCSGHA
		1	ŀ			NICHLHTGKCFCTTKGIKGDOC
					ŀ	OLCDSENRYVGNPLRGTCYYK
					ĺ	T\LIDYQFTF\SLLQEDDRHHTAI
		l			ļ	YFIVNPKQSNKNLHISINASNNF
					ŀ	NLNITWPC*STAGTISGEETYIV
						SKNNILEYRDSFSYEKFNFRSNP
					l	NITFYVYVSNFSWPIKIQIAFSQ
					1	HNTIMDLVOFFVTFFSCFLSLLL
		l				
		1			İ	EAAVVWKIKQTCWASRRREQL
		1				LRERQQMASRPFASVDVALEV
						GAEQTEFLRGPLEGAPKPIAIEP
						CAGNRAAVLTVFLCLPRGSSGA
						PPPGQSGLAIASALIDISQQKAS
00.581	50000	_	22522	2.52	***	DSKDKTSGVRNRKHLSTRQGT
22571	52939	В	22699	252	360	
22572	52940	A	22700	1	84	
22573	52941	A	22701	1	404	1000 to 00 000 000 000 000 000 000 000 00
22574	52942	A	22702	86	292	NPEMRCREFCLPSCRPELALE*
			1	1		G*SVRDVA*AQMCSVVWAGRP
						LGFAA*S*GSGHSCGRCSIWGE
		<u>_</u>		L		CWG
22575	52943	Α	22703	82	379	HTLQPFISMGSLTKMFKVISEGP
						MD*RPHPKVRPMT*HSTPAPAN
						FMGHFFWKIMICTDRTEVATLV
			1			AGV VSGASVGGKSPPYRKWNS
						PEPDVKGPRVL
22576	52944	Α	22704	2	382	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22577	52945	A	22705	1	479	HMPSGETAPLRIVDTGTGTGTV
						\AYAPTE\LGLHEMHIKYMASHI
						LESPLRFYVTTPTVPSVSAYGPG
						LVYGVAN/NTATFTIVTEDAGE
		1	ł			GGLDLAIEGPSKAEEPRPRLSRA
		1				PGVDGAGASPGLGEDTGPQAP
						GLQVLTERIPGPSAPTPEFLSVG VSAVC
22578	52946	A	22706	1	590	VSAVC
22579		A	22707	171	406	
	52947			269	2477	
22580	52948	A	22708		=	
22581	52949	В	22709	167	7472	
22582	52950	Α	22710	1	520	
22583	52951	Α	22711	2218	2472	
22584	52952	A	22712	722	1555	SRNWYEALNNTWGREDICVVD
ŀ						GWQSFRRQDYAFLQERGSGGD
		l				LWSSTLAQAMTLVESRHPRHQ
ļ		1			- 50	ALSTQLLDAIMPYCGNTLRLGV
						TGTPGAGKSTFLEAFGMLLIRE
ŀ		1				GLKVAVIAVDPSSPVTGGSILG
		1				DKTRMNDLARAEAAFIRPVPSS
			ĺ			GHLGGASQRARELMLLCEAAG
	i	1				YDVVIVETVGVGO\SETEVARM
			i			VDCFISLOIAGGADDLOGIKKG
	1	1				LMEVADLIVINKDDGDNHTNV
	1					AIARHM/YESALHILRRPPMHR
				1		VPRNPKVNLLPGTAMANLLS
22585	52953	В	22713	381	425	
22586	52954	A	22714	1	1185	
22587	52955	A	22715	1	3009	
22588	52956	A	22716	22	527	YEGIVMKPLIIVNPADCIGC\RT
		1				CE\VACVVAHPSEOELNADVFL
						PRLKVORLDSISAPVMCHOCEN
1	1	ı	1	I	1	APCVGACPVGALTMGEQVVQT
l	1	1	1	1	1	NSARCIGCOSCVSACPFGMITIO
l	ĺ					SLPGDTROOIVKCDLCEOR*EG
		l				PACVESCPTQALQLLIERELRRV
1		l		I		ROORIVVTGENPL
22589	52957	A	22717	187	325	INQUITY TO ENTE
22309	36731	<u></u>	22/1/	107	323	

SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequenee	codon for last amino acid of peptide scquence	Amino acid sequence (X=Unknown, *-Stop codon, /-possible nucleotide deletion, *=possible nucleotide insertion)
52958 52958	A	22718		2240	IMRSDPDTSVNEIETKLSALLGS ETTGEILFDLLCANGPEWNRFV TLEMKYGRIMLDTAKIIDEODV PTHILSKLTFTLRNHPEGVVMK NFEVLOPLOSULSGLPLWYSER ILQQINQLTHYEPVIGIMGKTGA GKSSLCNALFAGESPYSDVAA GKSSLCNALFAGESPYSDVAA CTRDELRERLQIGEHFMTIVDLP GVGESGVRDTEYAALYREQLP GVGESGVRDTEYAALYREQLP TYPOLIGENT TO THE STREET TO TH

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22591	52959	Α	22719	1	2301	MALTSGVHTGEKPYKCAQCEE
						AFSRKSELIIHQIIHTGEKPYECT
						ECGKTFSRKSQLIIHQRTHTGEK
	1					PYKCTKCGKSFCQQSHLIGHQR
		i i				IHTGEQPYLGLSVAERPKEYED
						DRALKMAAPCGNSSREQPCCFL
		1				NKASEDAERAKPPS*EPGRLRT
			0.0			P*VAQPH*VPHFMATWVRKHP
	1	l			ŀ	VHDRQFRLHGQIRELNWDVVR
		l				PEQFTGVATNCSGITYPQGGWL
		l				CPAELTRNVLELAQQQGLQIYY
	1		İ			QYQLQNLSRKDDCWLLNFAGD
						QQATHSVVVLANGHQISRFSQT
		1	1			STLPVYSVAGQVSHIPTTPELAE
						LKQVLCYDGYLTPQNPANQHH
1		1				CIGASYHRGSEDTAYSEDDQQQ
		l				NRQRLIDCFPQAQWAKEVDVS
			İ	l		DKEARCGVRCATRDHLPMVGN
				l		VPDYEATLVEYASLAEQKDEA
		l		l		VSAPVFDDLFMFAALGSRGLCS
				l		APLCAEILAAQMSDEPIPMDAS
			1			TLAALNPNRLWVRKLLKGKAV
22592	52960	A	22720	1021	1734	KAG
22593	52961	Ā	22721	1247	2412	RRSHHAANLDIRIRMPQTRQVT
	12201	ľ		1.2		LPDPELHPDSTLSMWPD/NRIAR
ļ		1				DAHYLYRYDRHGRLTEKTDLIP
						EGVIRTDDERTHRYHYDSOHRL
		l				VHYTRTQYEEPLVESRYLYDPL
ĺ		l				GRRVAKRVWRRERDLTGWMS
İ	1					LSRKPOVTWYGWDGDRLTTIQ
						NDRTRIQTIYOPGSFTPLIRVET
1	1					ATGELAKTORRSLADALQQSG
						GEDGGSVVFPPVLVOMLDRLE
	ļ			1		SEILADRVSEESRRWLASCGLT
						VEQMQNQMDPVYTPARKIHLY
						HCDHRGLPLALISKEGTTEWCA
						EYDEWGNLLNEENPHQLQQLI
					ĺ	RLPGQQYDEESGLYYNRHRYY
	1	1	l			DPLQGRYITQDPIGLKGGWNFY
			l	1		OYPLNPVTNTDPLGLEVFPRPFP
						LPIPWPKSPAQQQQWYPTFCC
22594	52962	Α	22722	1	4767	
-2007	1-2-02	۲.	1	1-	1	L

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22595	52963	Α	22723	668	1922	YPHADSPECWLSSSSERGKSSV
						TADLSRPRRLRPIPTCHGKPIAC
			İ			TTTKAKNAFRPIAGASATGRLA
						NSPIIILPNAAIRQEFGITLDPEEV
						FTFQGVKLYEYRYFPDCVRSRN
1					}	EAGILVEQILSTLFSVGEHFSLK
						RAPFFKVPSLRSFYAFGGLTTG
						VVRRNGNIIGVNTPARKIHLYH
						CDHRGLPLALISKEGTTEWCAE
						YDEWGNLLNEENPHQLQQLIR
	1	ı				LPGQQYDEESGLYYNRHRYYD
						PLQGRYITQDPIGLK\GGWNFY
	l					QYPLNPVQYIDSMGLASKYGH
	i					LNNGGYGARPNKPPTPDPSKLP
						DIAKQLRLPYPIDQASSAHNVF
			ŀ		İ	KTFFRALSPYDYTLYCRKWVK
1					Į.	PNLTCTPQDDSQYPGMDTKTA
						SDYLPQTN/WPVPSSIVSCTTGV
						VP*VVCQWVFAWPPDISHRRTG
22596	52964	Α	22724	25	510	RITDFGVKKIIRVGSCGAVLPHV
						KLRDV/VI/GMGAC/TDSKVNRI
						RFKGQAFGGIADFAMGRTAVD
						PGKALGINARVGNLVSADLFYS
	ļ					PDGEMFDVMEKYGILGVEMEA
	1					AGIYGVAAEFGAKALTICTVSD
	1					HIRTHEQTTAAERQTTFNDMIKI
						ALESVLLGDKE
22597	52965	A	22725	415	759	HRRRIVIDDTGIFRPRNTAQQLR
						QRPIAMSASGVVQIILEGHWRT
1				į		HSLRNGFHRGFRL*SSPKVGMQ
						HGSRQVDHRTQRALRFFRQPGI
						DFLRPVFTPRRQRLAATNHFSR
	<u> </u>					FVEH

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22598	52966	A	22726	sequence 2	2329	GTLPGCGAMHTCPSGPATHLV RRRIVRLPACVVTRANVPTPSR PVNATIGAFHSG*MASQGRRLS AQYPWSPPQYAVARPVPTVPCR LVPTGTSKAFKISVMOFOMGIP SCSISAKELITDFGVKHIRVGSG GAVLPHVKLRDVVIGMGACTD SKVTRIRFKDHAFAAIADFDMV RNAVDAAKALGIDARVGNLFS SADLFYSPDGEMFDVMEKYGIL GVEMEAAGIYGVAAEFGAKAI TICTVSDHIRTHEQTTAABSQT FINDMIKIALESVLLGDKEEDRV RFGKARATRYALLRPYRGIERII VRWYDDTGKAHKFADIRLCWI QGSCLDVAKAREVIRGSQPLPA WCENWAQGVDARLTKEAQSA LRPVINLTGTVLHTNLGRALQA EAAVEAVAQAMRSPVTLEYDL DAGGGHRDRALAQLLCRITG AEDACIVNNNAAVLLMLAAT ASGKEVVVSRGELVEIGGAFRI PDVMRQAGCTLHEVGTTNRTH ANDYRQADFQEDDFVMAFTGM INGPYQNTRALHGGHRRPDFLP
			20505		200	LLGAFHGAIDILAPGGLHGYSL PSSFFHALALSAAFPGKRCKLS VHLPFWGLKDGGPLLTAPLGG ALVGIVGGVGGSDLTFPFHTHS LNVRFVIRRISHRHHHFYRIAKL LMNGAQGSNMLIAAVAGVPHC NGHFPFTVADRFAIVLSEGSIRV
22599 22600	52967 52968	A	22727	876	702 1318	RCHLANASPAAGSHAPSPDSLS KIR/QEVG/PPLP/PLLAPLIATPP RTSQPLSPLISSSSPSSPASPVGQ VSPFRETPVPPAMSPWPEDPRR ASPPDPSPSPSAASASERVVPSP LQFCAATPKHALPVPGRLPPCA SGHAMPCWRAKIK

SEO ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		•
22.601	10000	_	22729	485	1455	PVVORARNASSCOPFSTSVWAI
22601	52969	Α	22129	483	1455	
		l			1	PSISATSVPMCGATHSILSPKKS
						TVSDRIGSMQISFFPLSRSAEK*
					1	ESPCSSDAFHAIFSVLSGLALHS
					i	TTTSLCSSTSGQLVCC**TSLPPT
					1	TYGMIACAAPVE*SPRCPV*PPA
						SAI*RCNNVEALCSTPFERQP*E
						PAK\SMPDHKPHECVYVVGE*D
						PAPHPNSREQIHPAHECLAGFP
		l			İ	VTQESLFAPLEIVPAYCCEPGRE
		l				PRSARSFPFAPRELT*VQLELGT
						GL*RETLDLVVSAGWA*ARRIS
		l				GA VTATPETGTLCQLDSGAECH
		l				YRETGVELP*RSPRPVGHGSSR
						HSRRWRT
22602	52970	A	22730	814	5001	PENSSHRPKPIIPMIASTLINANQ
						NSVSPYKRTFTRLTALMIIKKA
		1				AAQIHVGTSGNQYWRKLDFNS
		l				PGGSSPVENSDCSTNSRLSFSPE
	l	l				NILIQNQDIVREAAVQGDGQKQ
		l				RQPQATDLDSSGTHGSEMLPAT
		l				EVTVSGGFSVEETSCGDTGRSG
		l				GEALAVANDSTSTPQNANGLW
		l				KLKSTTPGGALPECFGTTDTTFS
		l				SAFCRKHGETQDTSQSSLPGTL
		l				HCYTGIREGGDDTEVESEAFSC
						SEGSEQQDAPD
22603	52971	Α	22731	34	102	
22604	52972	Α	22732	584	893	QKLLPNRMGHPVFVKSSQDPV
						LLSRIAWKDASTVTWVIEQDPV
		l				LRRRRRRTRRRRTRMRRRRGR*
		l				*/SKRRRMR/GRRGRR*CRRRRR
						RSSRSRPTRRQRNRNQV
22605	52973	Α	22733	2	269	IGKLYWFMPPSANLALRELGIV
			l			LFLSVVGLKSGLSWIGYGALIT
1		İ				AVPLITVGILARMLAKMNYLT
	l	l		,		MCGMLAGSMTDPPA*PRFQTN
1		l				DREEQHDPQLPQREVGA WRHK
		l				PVQLAD
			·		l	L`

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /-possible nucleotide
	sequence		09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22606	52974	A	22734	3	958	RCYSLLSRAPHPGIYISIAERV/*I
22000	32914	^	22734	ľ	/50	RLNSANFTLNSLLPAEKKLAEE
		1				FAVSRMTIRKAIDLLVAWGLV
						VRRHGSGTYLVRKDVLHQTAS
						LTGLGEVLKRQGKTVTSQVLIF
						EIMPAPPAIASQLRIQINEQIYFS
						RRVRFV\EGIPLMLEDSYMPVK
		1				LFRNLSLQHLEGSKFEYIEQECG
		1				ILIGGNYESLTPVLADRLLARQ
		1				MKVAEHTPLLRITSLSYSESGEF
						LNYSVMFRNATQIQGWRSHFR
					İ	R\LHPKKSLTVPPERAPPVFGGL
						IMRKNHSQRIDSGVRERRRTAG
		1				WMKIISKRORRRIGHGTCOHPA
		1				HGOVIHFG
22607	52975	Α	22735	2	285	GDEMTTSTLOKAIDLVTKATEE
		1				DKAKNYEEALRLYQHPVEYFL
					į.	HAVKYEAHSDKAKESIRAKCV
						QYLDRAEKLKDYLRSKEKHGK
						LCLKDRVT
22608	52976	A	22736	545	791	LCLRURVI
22609	52977	A	22737	249	1574	NHKAIDLVTKATEEDKAKNYE
22009	32711	^	22/3/	249	1374	EALRLYOHAVEYFLHAIKYEA
						HSDKAKESIRAKCVOYLYRAE
		1			ļ	KLKDYLRSKEKHGKKPVKENQ
		l				SEGKGSDSDSEGDNPEKKKLOE
					1	
						QLMGAVVMEKP\NIRWNDVAG
						LEGAKEALKEAVILPIKFPHLFT
	1					GKRTPWRGI\LLFRPPGT\GNSN
		l				RPKAGAPKATNSTFFSVSS\SDV
		ĺ	ŀ	l	l	KSKRLGETEKLV*NLF\ELARQ
				l	1	HK\PSIIFIDEVDSLCGSRNENES\
i			İ			QAARRIQNGSSLVQIAGGLGIN
ŀ				i		NDGDSGFFGATNIP\WVLD\SAI
	İ					R\RKV*KTKFYIPLPEKACPRPD
		i		l		VSRLHLGSTPHN\LRVATFHKL
				1		ARKTEGYSGADISIIVRDSLMQP
1		1	l	l		VRKVQSATHFKKVCGPSRTNPS
1		l		I		MMIDDLLTPCSPGDPRAMEMT
I		l	l	I	l	WMDVPGDKLLEPVVCMSDML
l		l	l	I		RSLATTRPTVNADDLL\KVKKF
						SEDFGQES
22610	52978	A	22738	52	567	
22611	52979	В	22739	1	1293	
22612	52980	Α	22740	443	1027	
22613	52981	A	22741	2077	2430	
22614	52982	Α	22742	1	504	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
					<u> </u>	
22615	52983	A	22743	3244	3947	RNEITPFRAMMIIRRMAHRFQT
	l	l		l		EEIKRISFIKRRVGNKDEINPGA
				1		RWRRFHDDGITFLRLLDAGLTI
		l				PSGKTIVHPLARQRRGLVKIQFL
			l		İ	R*LLAKLTRQPLLPLQPVTARL
			İ			CLNPRLRRQRKIRMPAGTKWQ
					l	RLCPLIGDR*RHRQRSAVYALS
		1				VRHNPVHHILRKNGDMYSFPIT
			i		1	IGNNVWIESHVVINPGVTIGDNS
		1				VIGAGSIVTKDIPPNVVAAGVP
						CRVILEINDRDK
22616	52984	Α	22744	742	1158	
22617	52985	A	22745	439	2882	PDWQAFQSPPFPASLIIHPKPAK
						LPGWLCIVQWSLLAITRTPTPV
						RWAOOTTETVGLVVGDVSDPF
		ŀ			ŀ	FGAMVKAVEQVAYHTGNFLLI
		1			ł	GNGYHNEQKERQAIEQLIRHRC
		1			ŀ	AALVVHAKMIPDADLASLMKO
		1				MPADKOCESIPDONNOTGTOY
		ı		ł	ł	PRGKEPKLLRRLMHMNRKVGP
		1				RGPRRMLLRYLDVEFLSVALA
		1				VVHLGPQGTCSRLFLVARALV
						VAVSEMDKSLSAQAQIYFASSN
		1				GPVPDLAYOVDFPRLEIGL*F\E
		l				YVDTGAETTLVPGDVLYVPAG
		1				GWNFPQWQAPATTFSVLFGKQ
		ı				QLGF/SAVVQWDGKQYQNLAK
		ŀ				OHVARRGPRIGSFLLOTLNEMO
		ŀ				MQPQEQQTARLIVASLLNHCR
						DLLGSQIQTASRSQALFEAIRDY
						IDERYASALTRESVAQAFYISPN
		ŀ				YLSHLFQKTGAIGFNENLSHTR LGHA*TLLKGYDLKVKEVAHA
		1	1	l	l	CGFVDSNYFCRLFRKNTERSPS
		1		l		EYRRHRRLHDNRICQFKTPTDN
	l	l				PENPDRYLTFRRIDKETSSLPKK
	1	l				VAPSCKRLSVSRSSPKPSISIRRC
		1				MRLMLIEGRRSSFIHPDKDVDG
		1				FHPYNVGRLCQRAPRLRPCTPR
		l		l		GIVTLLERYNIDTFGLNAVVIGA
		I		l		SNIVGRPMSMELLLAGCTTTVT
		l				HRFTKNLRHHVENADLLIVAV
		1	1	l		GKPGFIPGDWIKEGAIVMMSGI
		<u></u>				NRGVATDASCFPLVPFANRVSG
22618	52986	Α	22746	2	126	
22619	52987	A	22747	1	465	
22620	52988	С	22748	78	308	

SEQ ID	SEQ ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletinn, \-possible nucleotide insertion)
-				sequence		
	<u> </u>	<u> </u>				
22621	52989	Α	22749	I	2797	MEHPPVLAVQRVALPSRSSEPE
	i	ŀ				LFLNPPTDEETEAHRGMYLDQ
1	1					GPRTKDWIPVAPGVREPSPLGP
1						GPSSLWLLVVGGTVEEGSRKH
						RPTCKKDFTKSKTVDQAPTLCA
						RHQMLNIQFLISDRDPQCNLHC
					ŀ	SRTQPKPICASDGRSYESMCEY
		1				QRAKCRDPTLGVVHRGRCKDA
i						GQSKCRLERAQALEQAKKPQE
		1				AVFVPECGEDGSFTQNTSCIHSS
				İ		VYQIGENSSLTFLEKREESGHL
				!		QRDWFPLWFPRKARVI
22622	52990	Α	22750	1	5295	MNLNQEEIPDLPEKEFRRLVIKI
		1		l		IREGLEKGKAQCKEIQKMIQEV
1		1		i		KREIFQEIESLKKKPPKIQKTLDI
		l		l		LLKMONALESLSDRTEOAEEIN
		l				SELEDKVFKLTQSNKDKEKITR
1		i i				KYEOSLOEVWSYVKRPNLRIIG
		l	i			VPEEEENTKSLENILGGIIEENFP
l		l		ł	1	SFARDLDIQIPEAQRTTGKLIAK
		l				RSLLRHIVFRLSKVKMKERILR
		l		İ		AVROKHOATVSOKNKAGGITL
		1				PDFKLYYKATVSQKNKAGGITL
1		1				PDFKLYYK
22623	52991	Α	2275I	1	891	GSPFEVMPRDDVFFIVDVKI*DI
1		1				GVYSCTAONSAGSISANATLTV
		1				LETPSFLRPLLDRTVTKGETAV
		ı	1			LQCIAGGSPPPKLNWTKDDSPL
		1				VVTERHFFAAGNQLLIIGDSDV
1			ŀ	İ		SDAGKYTCEMSNTLGTERGNV
1						RLSVIPTPTCDSPQMTVL/HLLD
						DDGWGHCGLYVIIAVVCCVVG
	l	l	İ			TSLVWVVIIYHTRRRNEDCSITN
						TDETNLPADIPSYLSSOGTLADR
		l				ODGYVSSESGSHHOFVTSSGAG
						FFLPQHDSSGTCHIDNSSEADVE
						AATDLFLCPFLGSTGPYVFEGK
						CVWAONPF
22624	52992	A	22752	1	2947	MSAPSLRARAAGLGLLLCAVL
	1					GRAGRSDSGGRGELGOPSGVA
		1				AERPCPTTCRCLGDLLDCSRKR
		1		l		LARLPEPLPSWVARLDLSHNRL
		1				SFIKASSMSHLQSLREVAGNRIV
		1	1			EILPEHLKEFOSLETLDLSSNNIS
		1				ELOTAFPALOLKYLELNRNKIK
		l				NVDGLTFQGLGALKSLKMQRN
1				l		GVTKLMDGAFWGLSNMEILQL
1		1				DHNNLTEITKGWLYGLLMLQE
1		1	1			LHLSQNAINRISPDA WEFCQKL
		1				SELDLTFNHLSRLDD
22625	52993	A	22753	2	417	
22023	02//3	l.,	22177	<u>-</u>	1	L

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
22626	52994	A	22754	I	262	VLDIGS/SLTVPDEFSKQE/RAD GMWWKQLVAGAVAGAVS/RT GTAPLDRLKVFMQIKRAILGQQ ETLMCRSASWLAPWLVPQPNH HLPYG
22627	52995	Α	22755	1	762	
222628	52996	Α	22756		1527	ILLAAGADPAGSKGGGSREA PTEAARCRASPPRAGAMRGSP GDAERRQRWGRLFEELDSNKD GRVDVHELRQGLARLGGGNPD PGAQQGISSEGDADPNGGLDLE EFSRYLOEREGRLLLMFISLDR NQDGHIDVSEIQQSFRALGISIL LEQAEKNFAQA/VDRDGTMTID WQEWRDHFLLHSLENVEDVLY FWKHSTVLDIGECLTVPDEFSK QEKLTGMWWKQLAAGAVAG AVSRTGTVPLDRILKVFMQVHA SKTYNELNILGGI-SMYLEGGIRS LWRGNGINVLKIAPESAIKFMA GSLAGATAQTINYPMEVLKTRL TLRRITGQYKGLAGLRGGILER EGPRAFYRGYLPNVLGIIPYAGI DLAVYETILKNWWLQQYSHDS APGILVLACGTISSTCQJIAS YPLALVRTRMQAQSIEGGPKL SMLGLRHILSQEGMRGLYRGT SMYLERFYAGYLPNVLGIIPYAGI SMLGLRHILSQEGMRGLYRGT SMLGRHGWGYNGYMGYMGYMGYMGYNGYMGYNGYMGYNGYMGYNGYMGYNGYMGYNGYMGYNGYMGYMGYMGYMGYMGYMGYMGYMGYMGYMGYMGYMGYMG

SEO ID	SEQ ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		-
22629	52997	A	22757	1	2127	MDQSGLLALRAAVGAVVGSTL
						AVDIGILVKDWHPRPLGGKHC
						QLLGAFRCRQHRRGKEERRGE
						TKEKKGERENERGKRRRIEGER
						DKYVKENEKQTCVQKERHSCT
			İ			GNSPNHMATVQAERGALDSHG
						GWHPWDYCAVQQGRYHQLLL
						TKEEIEVQRSKKMCLLQAFLIC
						CQKSHFLGPGCTRSMVPASASG
						ESLKLLPLMVEDEGEPVTERE\R
					1	KREGERERGRERERRRRKRK
						RRRKRRRKRRRRRRRG*EFLS
						LG*NERGKRRRIEGERDKYVKE
		1				NEKQTCVQKERHSCTGNSPNH
		1				MATVQAERGALDSHGGWHPW
		1			1	DYCAVQQGRYHQLLLTKEEIE
1						VQRSKKMCLLQAFLICCQKSHF
		ļ.				LGPGCTRSMVPASASGESLKLL
		į .			1	PLMVEDEGEPVTEREQEEGGGE
				ł		REREREKKEKEKEKEKEEK
						EKEKEKEEERIRVPLPRVELRLT
1		ı			ì	NIAAKQVGNTILDVAAPADHPS
1						ESSCVYVRFVIKIPVAASFRFKK
		[SYVTNNVESCEDGIHYFWKSH
		1				MHSENSCSCKLPCPPHLPAAER
		ı				LGDALEPLLPTMRRQRQCQNT
		l	ļ	i		GDCDGRRQIGTRETEMHRDTSL
ł		l				VAVKTQVDTSREGMGEVEPNS
		1				CFWRDTHMASGVGAAFEELPH
1		1				DGTCDECEPDEAPGAEEVCREC
i		1				GFCYCRRHAEAHRQKFLSHHL
1		1				AEYVHGSQAWTPPADGEGAGK
22630	52998	A	22758	348	881	WROPRWPHHOACLWORAGSS
		1			ĺ	YSOHKDPHFTSFSTPRAVONFD
		l				GFPCPFLRETTTLVLIPSVPLTDP
l		ł				GSFYRAWSFQCGPRSPRPLLGR
İ		1				TLSQKFLLRLRGRPRRPYMSTL
ľ		1				TPF*STRIFERAFCQVMGAESGG
i						RIEKTTLCKRSLGPLMEPRV
22631	52999	A	22759	I	347	MVPASASGESLKLLPLMVEDER
				1	Ī · ·	EPVTERE/REKGRGREREGERE
				1	l	KEREGEREREREGERKGERER
				1		RRG*QPPFHNRFSEGQGRSFLCS
						RRPRVSRRGHTSRGPARVKAAE
						RRGLG
22632	53000	A	22760	145	258	PENGLIGLS*KIVLSLNIPVNEN
12032	155000	l.,	/-	1	1	DMKVDEVLYEDSSTA
L			<u> </u>	L	l	DMICTODIOLOGIA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22633	53001	Α	22761	1	1479	MALKEIFCPVRETETCTINCGTR
						ISPSILHVGLPYPSTPNVKPERE
		1				GCGGGVSGNSKQPGALETAESI
		1				GKKEGRREEGSRPESASSLSLLL
		1				SLAYDVDKSAQICTRIHSSVPLE
		1			1	AVYPGYIHQSIINQADLSALAPY
						DPCVCQIHKYHGAYNCLQYSV
		1				QNTLYSPTTEQYKRCQAPWVL
	Ì	1				PAIFTIPASGAAASAVPVRAQA
		1				QGGQAAGQRRRLATAPLTGEG
	1					EPDHGDRTRAGSQGPEARPRIH
			ļ			LGAARGGDRTRGRPGRRGRDW
		1				TRLGREMAAAAVSSAKRSLRG
						ELKQRLRAMSAEERLRQSRVLS
		1		ļ		QK\VIAHSEYQKSKRISIFLSMQ
		1				DEIETEEIIKD\ICQRGK\ICFIPPV
		1				RFSRAITMDMVRIGSP\EGNFFN
		ı				FPKTSWEYSLRLGEG\DVREEG
	1					LVPPGGL/EILILPCPQALGLDQ
		I				NHGQPRLG\RGQGAYL*CLS*K
1	į.	1				RLFCSHSGKLKPPTPPGGWLFK
						EQICPPRSPVNENDMK\VDEVL
22634	53002	A	22762	1	813	
22635	53003	Α	22763	1	368	
22636	53004	Α	22764	327	4281	SASASRCDPGSGSRREREEELQ
İ		1	ŀ			WRRRRRRRRRRRRRQRQRA
1		1				AAPAAPAGGIEAVNMASASYHI
Į.		1				SNLLEKMTSSDKDFRFMATND
						LMTELQKDSIKLDDDSERKVV
						KMILKLLEDKNGEVQNLAVKC
			l			LGPLVSKVKEYQVETIVDTLCT
						NMLSDKEQLRDISSIGLKTVIGE
		1	1			LPPASSGSALAANVCKKITGRL
			l			TSAIAKQEDVSVQLEALDIMAD
			ŀ			MLSRQGGLLVNFHPSILTCLLP
	ļ	Ļ				QLTSPRLAVRKRTIIA
22637	53005	A	22765	2	271	
22638	53006	A	22766	299	451	IVALFRSLVRHLRGTMAMESTA
				1	1	TAAVAAEL/VSADKIEDAPAPST
1	1	1	I			SADKVES

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22639	53007	A	22767	1	1122	MATESTATAAIAAELVSANKIE
22039	33007	<u> </u> ^	22707	ľ	1122	DVPAPSTSADKVERIENGVLGN
			İ			TLDGVHVEEEEGEKTEDESLVE
						NNDSIDEEARKELREOVYDSM
			l	i		GEKEEAKKTEDKSFAKPETDKE
						ODCEMEKGGREDMDISESAEEP
	ŀ		İ			OEKVDLTLGWLTEISEGAKGG
						GAPEGPNEAEVTSGKPEOEVPD
						VEEEKSVSETDVQKECREKGG
						WEKHREVIVSIEEKPKEVSEEOP
						VVILEKQGTAVEVAAESLDPTV
l						KPVDVGGDEPEEK/GTVPGSLR
		1		ł		
1		1			1	PSPCRDPLPAGLGLWVQLSV**
				ł		GSGTVQQIY*SH*EENGCTKRA
		1				GEGG*RIVC*IQERN*GTEGTAT
		1		1		RNYRKDRRCKGVSDYWECS*T
22640	50000	١.	007/0	222	339	GSESYSGGELYFRFHS*WRRLF
22640	53008	A	22768	223	496	
22641	53009	В	22769	12	1099	VSGVPNCLFFSOALFRSLVRHL
22642	53010	A	22770	12	1099	
		1				RGTMAMESTATAAVAAELVSA
			1			DKIEDVPAP\STS\ADKVESLDV
		1			ŀ	DSEAKKLLGLGQK\HLVMGDIP
				l	İ	AAVNAFQEAASLLGKKYGETA
		1				NECGEAFFFYG\KSLLELARME
		l			l	NGVLGNA\LEGVHVEEEEGEKT
1		l				EDESLVENNDNI/D*N*RLGRG*
1						*RK**D*RNAK*FSP*KQVSSRK
		1				*GGGDWEPRACLGYAGFSKDH
Ì		1				F*KARNKRSTALCCPGTS*TRRS
					1	*C*I*\KLCASC\GGVPVLP*PTGP
		1	1	i	1	VPGSPRPSPCRDPLPAGLGLWV
		l				QLSV**GSGTVQQIY*SH*EQNG
		l			l	CTKRAGEGG*RIVC*IQERN*GT
		İ				KGTATRN*REDRRCKGVSA*W
1		l		L		ECS*TGSESYS

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
		<u> </u>			Ļ	
22643	53011	A	22771	11	2485	VSGVPNCLFFSQALFRSLVRHL
						RGTMAMESTATAAVAADVVS
						ADKIEDVPAPSTSADKVESLDV
						DSEAKKLLGLGQKHLVMGGIP
			l			AAV\NAFQGAA\SLL\GKKYGET
						ANE\CGEAFFFYG\KSLLELARM
	i		İ			ENGVLGNALEGVHVEEEEG\EK
	1					TEDESLVENNDN\IDEEAREELR
						EQVYDAMGEKEEAKKTEDKSL
	1					AKPETDKEODSEMEKGGREDM
1						DISKSAEEPQEKVDLTLDWLTE
						TSEEAKGGAAPEGPNEAEVTSG
	1					KPGQ\EVPDAEEEKSVSGTDVQ
						EECREKGGOEKOGEVIVSIEEK
						PKEVSEEOPVVTLEKOGTAVEV
					ľ	EAESLDPTVKPVDVGGDEPEEK
						VVTSENEAGKAVLEOLVGOEV
	1					PPAEESPEVQTEAAEASAVEAG
						SEVSEKPGQEAPVLPKDGAVN
		l				GPSVVGDQTPIEPQTSIERLTET
	i					KDGSGLEEKVRAKLVPSQEETK
	İ		l			LSVEESEAAGDGVDTKVAQGA
						TEKSPEDKVQIAANEETQEREE
1						QMKEGEETEGSEEDDKENDKT
1						EEMPNDSVLENKSLQENEEEEI
						GNLELA WDMLDLAKIIFKRQET
		l	i .			KEAQLYAAQAHLKLGEVSVES
	ĺ					ENYVQAVEEFQSCLNLQEQYL
			İ			EAHDRLLAETHYQLGLAYGYN
						SQYDEAVAQFSKSIEVIENRMA
				i		VLNEQVKEAEGSSAEYKKEIEE
				1		LKELLP\EIREKIENAKE\SQR\SG
22644	53012	Λ	22772	52	161	FONPTGLY/CDTEYN\RLOKOO
			_			TEKTKOKKTOFHPICN
22645	53013	A	22773	3	115	FFQNPTGLY/CDTEYN/RLQKQQ
		1		1		TEKTKOKKTOFHPICN
22646	53014	A	22774	229	358	
22647	53015	A	22775	257	386	
22648	53016	A	22776	259	294	
22649	53017	A	22777	217	398	KAWWNLLRKSEPKSCGS*GGL
22049	33017	l^	22///	1217	370	STSRERLVECSSPLGLGYCG**L
1	1	1	l	l		CHLOEPHYGSLHRMSS
22650	52010		22770	l	2204	CHEQUETT I GSETIKMSS
22650	53018	A	22778	11	3386	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22651	53019	A	22779	391	2100	SRRTAGVRLDILCKDPIMWVSA
						DGHSYQPQYPKAKRATEFHFF
						NFKALLARAAVGAARGIHIHRC
		l				GLTKRPDARRRRVAGAOGREP
				i		AISLPGDRAAGARATRTRGPGP
		ĺ				APKMPAIAVLAAAAAA\WCFL
1						QVESRHLDALAGGAGPNHG\NF
						L\DNDQ\WLSTVSQYDRDKY\W
						NRFRDEVEDDYFRNW\NPNKPF
						DQAL\DPSKDPCLKVKCSPHKV\
						CVTQ\DYQTAL\CVSRKHLLPR
						QKEWGTWAQKHWGLDL*ILV
						K\CKPCSRGRSSAMGLAGSDGP
	l					LLTAFQVANLEFHACSTGQK/S
	ŀ		į.			FATLCDG\PCPCSSQSLEPPKHK
					ļ	GRKGVPCTDKELRNLASRLKD
						WFGALHEDANRVIKPTSSNTAQ
	İ				1	GRFDTSILPICKDSLGWMFNKL
					i	DMNYDLLLDPSEINAIYLDKYE
			•			PCIKPLFNSCDSFKDGKPFLNNE
						WCLLPSQNPGGLP/CAQNEMNR
		1				IQ\KLSKGKSLLGAFIPRCNEEG
		l			ľ	YYKATQCHGSTGQCWCVDKY
1						GNELAGSRKQGAVSCEEEQETS
1						GDFGSGGSVVLLDDLEYERELG
						PKDKEGKLRVHTRAVTEDDED
						EDDDKEDEVGYIW
22652	53020	A	22780	2	441	HGVQGAGCGEVAAVSIELGAG
						GPVWRAAPRACSRLQLLRGFG
						LEGQWFGESCCRRGAS*EGSTR
						*PPPRRQPWPEWVMSWVPPET
				ŀ		REPARISPAPAKQN*PPPAPKSP
1						AAPA\PQAAVAATPSPPGEISEA
		_				WDSLESFSKPAFFSQ
22653	53021	Α	22781	3	121	

SEQ ID	SEQ ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hed	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22654	53022	Α	22782	2	1488	FVNNMADGEEPEKKRRRIEELL
					ľ	AEKMAVDGGCGDTETWKDR\
						WNHVKK\FLERSGPFTHPDFEP\
						STESLQFLLDTCK\VLVIGAGG\
						LGCELLKNLALS\GFRQIHVID\
						MDTI\DV\$N\$*IRQFLFRPKDIGR
						PKAEVAAEFLNDRV\PNC\NVV
		l				PHF\NK\IQDFN\DTFY\RQFHI\IV
						CGTGTLSCARRWINGMLISL\LN
			l			YERMVS*DPSSIVPLID\GGTEGF
						\KENARVIL\PGMTACIECTLEL
						YPPQVNFPMCTIASMPRLPEHCI
						EYVGMLQWPKEQPFGEG\VPL
						DGDDP\EHIQWIFQKSLERASQY
						NIRGVTYRLTQGVVKRIIPAVA
						STNAVIAAVCAT*/EVFKIATSA
			i			YIPL*IITWVFNDV\DGLYTYTFE
						GRKGK\ENCPAC\SQLPQNISVF
						LHQAKLQ\EVLGIILTNSASLQN
		i				EILPAITATLGGEKIGTLYLQSV
						TSI\EERNRANFFQTIERNWGLL
1		1				DGPKTGRLLDVTTP\QTVLFQTS
				1		ILLLKGKSPT
22655	53023	A	22783	1	1719	
		_				
22656	53024	В	22784	199	460	
22656	53024 53025	B A	22784 22785	199 226	460 1620	NGSGLALATGNRDPGLCPLSAP
		_				SCPWR*KSAGASTQQAYSSQSQ
		_				SCPWR*KSAGASTQQAYSSQSQ /HPAYHRITPT\LTNFALRLYKEL
		_				SCPWR*KSAGASTQQAYSSQSQ /HPAYHRITPT\LTNFALRLYKEL AADAPGNIFFSPVSIS\TTLCLLC
		_				SCPWR*KSAGASTQQAYSSQSQ /HPAYHRITPT\LTNFALRLYKEL AADAPGNIFFSPVSIS\TTLCLLC LG\AQANNSALILEGLGFNLTET
		_				SCPWR*KSAGASTQQAYSSQSQ /HPAYHRITPTLTNFALRLYKEL AADAPGNIFFSPVSIS\TTLCLLC LG\AQANNSALILEGLGFNLTET PEADIHQGFRSLLHTLVLPSPKL
		_				SCPWR*KSAGASTQQAYSSQSQ HPAYHRITPTLTNFALRLYKEL AADAPGNIFFSPVSISTTLCLLC LGVAQANNSALILEGLGFNLTET PEADIHQGFRSLLHTLVLPSPKL ELKVGNSLFLDKRLKPRQHYFD
		_				SCPWR*KSAGASTQQAYSSQSQ #PAYHRITPTLTNFALRLYKEL AADAPGINFSPVSISTTLCLLC LGVAQANNSALILEGLGFNLTET PEADIHQGFRSLLHTLVLPSPKL ELKVGNSLFILDKRLKPRQHYFD SIKELYRAFAFSAHFTDSVTTGR
		_				SCPWR*KSAGASTQQAYSSQSQ /HPAYHRITPTLTNFALRLYKEL AADAPGNIFFSPVSIS\TTLCLLC LGYAQANNSALILEGLGFNLTET PEADIHQGFRSLLHTLVLPSPKL ELKVGNSLFLDKRLKPRQHYFD SIKELYRAFASAHFTDSVTTGR QINDYL*RRTYWPGVDLPFRS/L
		_				SCPWR*KSAGASTQQAYSSQSQ HPAYHRITPTLINFALRLYKEL AADAPONIFFSPVSISITILCLLC LGWQANNSALILEGLGFNLTET PEADIHQGFRSLLHTLVLPSPKL ELKVGNSLFLDKBLKFRQHYFD SIKEL YRAFAFSAHFTDSVTTGR QINDYL*RRTYWFGVDLPFRS/L ARDTFMVLANFILESKAKVKA
		_				SCPWR*KSAGASTQQAYSSQSQ HPAYHRITPTLTNFALRLYKEL AADAPONIFSPVSISITTLCLLC LG\(\)QANNSALILEGLGFNLTET PEADIHQGFRSILHTLVLPSPKL ELK\(\)GNSILHTLVLPSPKL ELK\(\)GNSILDKRLFRQHYFD SIKELYRAFAFSAHFTDSVTTGR QINDVL*RRTYWFGVOLPFRS/L ARDTFMVLAINFILFSKAK\(\)KA TLSSRVPDPKSRPVFLVDERTSF
		_				SCPWR*KSAGASTQQAYSSQSQ HPAYHRITPTLTNFALRLYKEL AADAPONIFFSPVSISITTLCLLC LGVQANNSALILEGLGFNLTET PEADIHQCFSSLLHTI-VLSPKL ELKVGNSLFILDKRLKPRQHYFD SIKELYRAFAFSAHFTDSVTTGR QINDVL*RRTYWFGVDLPFRS/L ARDTFMVLAINFILESKAKVKA TLSSRVPDPKSRPVFLVDERTSF QVPMMHQKEMHRFLYQQDLA
		_				SCPWR*KSAGASTQQAYSSQSQ HPAYHRITPTLTNFALRLYKEL AADAPGNIFFSPVSISITTLCLLC LGYQANNSALILEGLGFNLTET PEADIHQGFRSLLHTLVLPSPKL ELKVGNSLFLDKRLKFRQHYFD SIKEL YRAFAFSAHFTDSVTTGR QINDYL*RRTYWFGVDLPFRS/L ARDTFMVLANFILESKAKVKA TLSSRVPDPKSRPVFLVDERTSR QVPMMHQKEMHRFLYDQLLA CTVLQIESYRGNALALLVLPDFG CTVLQIESYRGNALALLVLPDFG
		_				SCPWR*KSAGASTQQAYSSQSQ #PAYHRITPTLTNFALRLYKEL AADAPONIFFSPVSISTTLCLLC LGXQANNSALILEGLGFNLTET FEADIHQGFSSLLHTLVL.PSPKL ELK-VGNSLFLDKRLK-PRQHYFD SIKEL YRAFAFSAHFTDSVTTGR QINDYL-PKRTYWFGVDLPFRS/L ARDITFMVLANFILFSKAKVKA TLSSRVPDPKSRPVELVDET LOYPMMHQKEMIRFLYDQDLA CTVLQIELYRGNALALLVL.PDG KMKQVEAALQPQTLRKWGQL
		_				SCPWR*KSAGASTQQAYSSQSQ HPAYHRITPTLTNFALRLYKEL LGAQANNSALILEGLGFNLTET PEADIHQGFRSLLHTLVLPSPKL ELKVGNSLFLDKRLKPRQHYFD SIKELYRAFAFSAHFTDSVTTGR GINDVL*RRTYWGVOLDFRS/L ARDTFMVLAINFILFSKAKVKA TLSSRVDPKSRPVLDERTSF QVPMMHQKEMHRFLYQOLLA CTVLQIEVRGHALLVLPPG KMKQVEAALQPQTLRKWGQL LLPSLLDLHPRFSISGTYNLEDI
		_				SCPWR*KSAGASTQQAYSSQSQ HPAYHRITPTLTNFALRLYKEL AADAPONIFFSPVSISITTLCLLC LGQAQANNSALILEGLGFNLTET: ELKVGNSLFILDKRLKPRCHYEP ELKVGNSLFILDKRLKPRCHYEP ELKVGNSLFILDKRLKPRCHYEP GVINDYL*RRTYWPGVDLPFRS/L ARDITMVLANIFILESKAKVKA TLSSRVPPRSRPVFLVDETST QVPMMHQKEMHRFLYDQDLA CTVLQIEVTRGMALALLYLDPE KMKQVEAALQPCITLRKWGQL LLPSLLDLHLPRSISGTYNLEDI LEPSNWFSPTYTLNKITSLQSVG
		_				SCPWR*KSAGASTQQAYSSQSQ #IPAYHRITPTLTNFALRLYKEL AADAPONIFFSPVSISITTLCLLC LGQQANNSALILEGLGFNLTET FEADIHQCFSSLLHTI.VL.SPSKL ELKVGNSLFLDKRLKPRQHYFD SIKELYRAFAFSAHFTDSVTTGR GINDYL*RRTYWFGVDLPFRS/L ARDTFMVLANFILESKAKVKA TLSSRVPDPSSRPVELVDETT QVPMMHQKEMHRFLYDQDLA CTVLQIEVRGNALALLVLPDFG KMQVPEAALQPOTLRKWGQL LLPSLLDLHLPRFSISGTYNLED LSPNWFSPTYTLNKITSLQSVG G*LNKSISKVSHKSMVDMMEK
		_				SCPWR*KSAGASTQQAYSSQSQ APPAYHRITPTLINFALRLYKEL LGAQANNSALILEGLGFNLTET PEADIHQGFRSILHTIVLPSPKL ELKVGNSI-FLDKRLKPRQHYFD SIKELYRAFAFSAHFTDSVTTGR GINDVL*RRTYWGFOULPFRS/L ARDTFMVLA\NFILFSKAKVKA TLSSRYDPKSRPVFLVDERTSF OVPMMHQKEMHRFLYDQDLA CTV-LQIEVYRGNALALLVLPDPG KMKQVEAALQPQTLTRKWGQL LLPSLLDLHPRSISGTYNLEDI LSPNWFSFTYTLNKITSLQSVG G*LNKSISKVSHKSMVDMNEK GDQRPGLLQASSPSPHL*YTMS
		_				SCPWR*KSAGASTQQAYSSQSQ #IPAYHRITPTLTNFALRLYKEL AADAPONIFFSPVSISTTLCLLC LGXQANNSALILEGLGFNLTET EEADHQGFRSELHTI-UL-PSPKL ELK-VGNSLF-LDKRLK-PR-QHYPE SIKEL YRAFAFSAHFTDSVTTGR QINDYL-PKRTYWFGVDL-PFSXL ARDITFMVLANIFILEFSKAK-VKA TLSSR-VPDFSSR-PVEL-VDET-SK- LOY-PMM+HQKEMHRFL-YD-QDL-A CTVLQIE-YRGNALALL-VL-PDFG KMKQVEAALQPQTL-RK-WGQL LLPSLLDLH-PFFSISGTYNLED LESPNWFSSTYTLNKIFISLOG G*LNKSISK-VSHKSMYDMNEK GDQRPGLLQASSPPL*-YTMS DPHAHFINRPFLLLL-WEVTTQSL
		_				SCPWR*KSAGASTQQAYSSQSQ APPAYHRITPTLINFALRLYKEL LGAQANNSALILEGLGFNLTET PEADIHQGFRSILHTIVLPSPKL ELKVGNSI-FLDKRLKPRQHYFD SIKELYRAFAFSAHFTDSVTTGR GINDVL*RRTYWGFOULPFRS/L ARDTFMVLA\NFILFSKAKVKA TLSSRYDPKSRPVFLVDERTSF OVPMMHQKEMHRFLYDQDLA CTV-LQIEVYRGNALALLVLPDPG KMKQVEAALQPQTLTRKWGQL LLPSLLDLHPRSISGTYNLEDI LSPNWFSFTYTLNKITSLQSVG G*LNKSISKVSHKSMVDMNEK GDQRPGLLQASSPSPHL*YTMS
		_	22785	226	1620	SCPWR*KSAGASTQQAYSSQSQ #IPAYHRITPTLTNFALRLYKEL AADAPONIFFSPVSISTTLCLLC LGXQANNSALILEGLGFNLTET EEADHQGFRSELHTI-UL-PSPKL ELK-VGNSLF-LDKRLK-PR-QHYPE SIKEL YRAFAFSAHFTDSVTTGR QINDYL-PKRTYWFGVDL-PFSXL ARDITFMVLANIFILEFSKAK-VKA TLSSR-VPDFSSR-PVEL-VDET-SK- LOY-PMM+HQKEMHRFL-YD-QDL-A CTVLQIE-YRGNALALL-VL-PDFG KMKQVEAALQPQTL-RK-WGQL LLPSLLDLH-PFFSISGTYNLED LESPNWFSSTYTLNKIFISLOG G*LNKSISK-VSHKSMYDMNEK GDQRPGLLQASSPPL*-YTMS DPHAHFINRPFLLLL-WEVTTQSL
		_				SCPWR*KSAGASTQQAYSSQSQ HPAYHRITPTLTNFALRLYKEL LGAQANNSALILEGLGFNLTET ERADHQCPSSELLHTI.VLSPKL ELKVGNSLFLDKRLKPRQHYFD SIKELYRAFAFSAHFTDSVTTGK GINDTL*RRTYWFGVDLPFRS/L ARDTFMVLAINFILESKAKVKA TLSSRVPDPKSRPVFLVDERTSF QVPMMHQKEMHRFLYDQDLA CTVLQIEVRGNALALLVLPDPG KMKQVPAALQPQTLRKWGQL LLPSLDLHLPRFSISGTYNLEDI LSPNWFSFTYTLNKITSLQSVG G*LNSISKYSKHSMVDMMEK GDQRPGLLQASSPSPHL*YYTMS DPHAHFNRPFLLLLWEVTTQSL LFLGKVVNPVVWTMVGQQE

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22660	53028	A	22788	309	2286	CAVCHISFQDTPVLVSSNVTMQ
		l				FGSKPLFENISVKFGGGNRYGL\
						IGANGSGKSTFMKILGGDLEPT
		1				LGNGSLDPNER\IGKLRQ\DQFA
		1				FEEFTVLDTVIMGHKEL\WKVK
	ŀ					QERDRIYALPEMSEEDGYKVA
	1	1				DLEVKYGEMAGYSAEARAGEL
					1	LLG\VGIPVEQHYGPMSEVAPG
	ł	İ			l	WKLRVLLAQALFADPDILLLDE
						PTNNLDIDTIRWLEQVLNENNS
		1				T\MIIISHDRHFLNMVCTHMAD
						LDYGELRVYPGNYDEYMTAAT
					1	QARERLLADNAKKKAQIAELQ
		l				SFIDKIKLEEVKASSRQNPFIRFE
						QDKKLFRNALEVEGLTKGFDN
	1				ł	GPLFKNLNLLLEVGEKLAVLG'T
						NGVGKSTL\LKTLVGNLQPDSG
	1				l .	TVKWSENARIGYYAQDHEYEF
						ENDLPVFEWMSQCKQ*GDDEQ
						AVRSIFGRLLFSQDVIKKPAKVL
						SAPFCKDSILEAIDAGIKLIITITE
	1				1	GIPTLDMLTVKVKLDEAGVRMI
	1			l	l	GPNCPGVITPGECKIGIQPGHIH
	1					KPGTGELNITTGGIVKARDTQIA
	1			ļ		LNDKSKGDVRVDGQNSLLETF
						NMYVGTSGTGTLTLTNNGTLN
	1	l				VEGGEVYLGVFEPAVGTLNIGA
	i	ŀ				AHGEAAADAGFITNATKVEFG
	1	l				LGEGVFVFNHTNNSDAGYQVD
						MLITGDDKDGKVIHDARHTVF
22661	53029	Α	22789	I	553	
22662	53030	Α	22790	1633	2207	ILFFDKGMPAQTPSLQKRWEK
	i	1				M*LVLW*GDREGIKAI/ISEESW
		l	l			GARAPIFLLR/MMEDQGLYSIR
		1				MGGARGLPGKSLAL*GAGRGF
						YSGERTWTLSLPSGNCAP\YQR
						GGWWYHACA\HSNLNGVWHH
						GGHYRSRYQDGVYWAEFRGG
				l		AYSLRKAAMPHLAPEAVTLCSF
		_				GPLGPRGHWSAGAPSCSGHTFF
22663	53031	Α	22791	34	210	AAADPLRHLRRGAFIKAGRHA
			l			G/PSLPVCLQATA\H*MDLARLA
	<u> </u>					LVLVVRTSASTKSSTRE

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22664	53032	l la	22792	397	1168	LLSRISKVSSKLPVRGGAGSLCT
						GDQRK\RRKFL\ETVELQISLKN
						YDPOEGOSAFSGHPSRL*V\PSP
		l				PLSSLVCVLGGPSKHL*RRLRP
		1				VGYPPTMDIEGA*KKLNOELKN
		İ				LVKKAGPRKYDCVFWPQESLD
		ı				SSKIPPNPSGPGFK*RAGKVPFP
		l				CLTHNGKHGLAQSGIEVEVPTF.
		l				LSFQMKKGVYVWAVAVWSRE
		l				R*QTNELVYNIHLACQLSWVSL
		1				LKEKNWQ\NVPGPYIIKSTMGQ
						SPSSHILRHIRINSITS
22665	53033	Α	22793	1	377	SVSRVPTRSLRTSTWRLAQDQT
		İ				QDTQLITVDEKLDITTLTGVPEE
		l				HIKTRKVRIFVPARNNMQSGVN
		l				NTKKWKMEFDTRERWENPLM
		l				GWAST\WSYDIEERKVPKPKSK
22555	50004	Ļ.	00004	100	260	SYGANFSWNKRTRVSTK
22666 22667	53034	A	22794	183	368 623	ILAFACSKMAAVSMSV\VLROT
22007	33033	^	22193	ľ	023	LWRRRAVAVAAL\SVSRVPT/R
		l		1		RSLRTSTWR\LAH\DQTQDTQLI
		l		i		TS**KIGISLTFNWPFPEEHIKNL
		l	i	l		EKSGIFVPCSPITCOSGVNNTK\
		l		l		KWK\MEFDT\RER\WEN\PLMG\
				ŀ		WASTGLIPLS\NMVLNLPVLKE
		ł		l		DAVS\FAEK\NGWELWTLEERK\
		ł				VPKPKSQVFMGANFS\WNKRTR
		1		1		VSTKIGLALTISPA
22668	53036	Α	22796	1	798	MNKVNPTQQDKEDHPELIITSIG
		l		1		DKRQLTLEAASGVCHRGTQTR
		1				VKEESKKFLISMSFRGGKPVHK
		l			İ	VVERACVCTCLHHTSVPATHN
		l		l		TQRRHTTPTTRKNGRKRRGKK
		l		l		EREKRKDREHNTGRRTKRKKT
		l				RNIGGQRTQQKQKQSEDKARR
				l		EPGGGRKQGRGREDGTRQRRG
		1	l			GRTRNREEGAQRGRNGR/RGEG
		1	l	l		RRTEEETHMA*APDITATQQTE
			l			QQGHGTRQRDTTLRRHTAEDA
						QPSEHASADQAHTKKTE*SVSR
22669	53037	В	22797	85	322	TNGQYLLT
22009	22027	ľ	44171	102	322	

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codnn for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			}	sequence		
22670	53038	A	22798	165	4733	AATALLTHGQEEGQVEGQDED
		1				IPPITCVQNGLRYHDRDVWKPE
						PCRICVCDNGKVLCDDVICDET
		1	i	1		KNCPGAEVPEGECCPVCPDGSR
		1		l		VTHRPKKPPGVEGPKGDTGPPR
		1				PKGT\PGPPGRDGIPGQPGLPGP
		i		l	ŀ	PGPPGPPGPPGLGGKLLLPKLSY
		i			-	GYDEKSTGGISVPGPMGPSGPR
	l			1		GLPGPPGAPVSIQDVFICLLGLW
	1				ŀ	SFGGKTGMRAGEMLRDLLGPO
				l	ŀ	GFQGPPGEPGEPGASGPMGPRG
	1			l	İ	PPGPPGKNGDDG
22671	53039	С	22799	88	392	PFGFFGKNGDDG
22672	53040	Ā	22800	37	293	
22673	53041	A	22801	55	1917	GGGGVRRRRGTRRSEPGDSRSA
22075	15501.	ľ.		-	1	MAPIGLKAVVGEKIMHDVIKK
						VKKKGEWKVLVVDQLSMRML
						SSCCKMTDIMTEGITIVEDINKR
				i	!	REPLPSLEAVYLITPSEKSVHSLI
					i	SDFKDPPTAKYRAAHVFFTDSC
						PDALFNELVKSRAAKVIKTLTEI
	l	1				
	l	l		1		NIAFLPYESQVYSLDSADSFQSF
		l		İ		YSPHKAQMKNPILERLAEQIAT
	l					LCAT\LKEYPA\VRYRGEYKDN
	l	l				ALLAQLIQDKLDAYKADDPTM
	l	1		·		GEGPDKARSQLLILDRGFDPSSP
			l			VLP*I*LFQGMSY*SCLPI\ENDV\
				1		YKYETSGIGGGHGVKGGASGT
	I	1			1	EDDDLW\IALR\HKHIAEV\SQ\E
	1	1				VTRSLKDFSSSKRMNTGEKTT
	İ		ĺ			MRDLSQMLKKMPQYQKELSK
	1					YSTHLHLAEDCMKHYQGTVDK
						LCRVEQDLAMGTDAEGEKIKD
	1		l			PMRAIVPILLDANVSTYDKIRIIL
1						LYIFLKNGITEENLNKLIQHAQI
	1					PPEDSEIITNMAHLGVPIVTDST
	1					LRRRSKPERKERISEQTYQLSR
						WTPIIKDIMEDTIEDKLDNK\HY
				ĺ		P\YISTR\$\$A\$F\$TTAV\\$ARYGH
	1	l		l		WP*GTRPPGEYRSGP/RVFIIFHP
						LGGVSLNEMRCALRR*PRANG
		l				KWGGA**GSTHILOPHRNLPGH
					L	E GOTT GOTTING OIT

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last eodon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, +=Stop codon, /-possible nucleotide deletion, \=possible nucleotide insertion)
22674	53042	A	22802	46	1931	GGGGVRMLCGTRRSEPGDSRS
						AMAPIGLIAVVGEKIMHDVMI.
	i					NDLNMGEWKALLVDQLSMRM
!	1					LSYCCLMTDIMTEGITIVEDINK
		l				RR*PLPSLEAVYLITPSEKSVHS
						LISDFKDPPTAKYRAAHVFFTD
						SCPDALFNELVKSRAAKVIKTL
	l					TEINIAFLPYESRVYSLDSADSF
	ŀ					QSFYSPHKAQMKNPILERLAEQ
						IATLCATLKEYPAVRYRGEYKD
	l					NALLAQLIQDKLDAYKADDPT
						MGEGPDKARSQLLILDRGFDPS
	l					SPVLHELTFQAMSYDLLPIEND
	ŀ					VYKYETSGIGEARVKEVLLDED
i	i					DDLWIALRHKHIAEVSQEVTRS
		1				LKDFSSSKRMNTGEKTTMRDL
						SQMLKKMPQYQKELSKYSTHL
						APCLRTCMKHYQGTVDKL\CR
						V\EQDLAMG\TDAEG\EKI\KDP
ł						MRAIVPILLDANVQTY*QIRIILL
ŀ						YIFLKNGITEENLNKLIQHAQIP
						PEDSEIITNMAHLGVPIVTDSTL
				1		RRRSKPERKERISEQTYQLSRW
		i				TPIIKDIMEDTIEDKLDTKHYPYI
	1	1			1	STRSSASFSTTAVSARYGHWHK
		1				NKAPGEYRSGPRLIIFILGGVSL
		l				NEMRCAYEVTQANGKWEVLIG
l	1	l			1	STHILTPTKFLMDLRHPDFRESS
						RVSFEDQAPTME
22675	53043	Α	22803	2	51	
22676	53044	A	22804	236	451	LHLPLFLGAIRSGQYGLWNLFR
		1				VLGWTCSLGIYLGWRVS*NFG
						RGTKALKVLSTIS/E*RNHHCHP *NNIFVY

SEQ ID NO:	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence	liou	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22677	53045	A	22805	3	1173	GAGAPAAGQAGLTSTSAVRAR
		l				PVSPARQSQDPAASLSLHLSLPP
l	1					APGAGCARWRRRGPQAAMAA
						GAGAGSAPRWLRALSEPLSAA
						QLRRLEEHRYSAAGVSLLEPPL
						QLYWTWLLQWIPLWMAPNSIT
	i					LLGLAVNVVTTLVLISYCLTAT
						EEAPYWTYLLCALGLFIYQPLD
		ļ				AIDGKQARRTNSCSPLGELFDH
		1		l .		GCDSLSTVFMAVGASIAARLGT
		i i				YPDWFFSRSFYLGCLVFYC\AH
		l	l			WQTYVS\GMLRFGKVDVTEIQI
		l				AL\VIVFVL\SAFG\GATM/WGD
	1	l				YTIPILEIKLKILPVLGFLGGVIFS
		l		l		CS\NYSHVILH\GGVGKNG\STIA
		l		l		GTSVLSPGTPH/MGLIIIL\AIMIY
		l				*KVRQLMVF*KSIPCL*YPQCLG
						CVF\AKVSQKISGSSHDQK
22678	53046	A	22806	2	1091	CPGRRRRRRPGRSRARATALPT
ļ		1				PGPRKRGGPAGRKRERPERCSS
		1				SSGGSSSSGDEDGLELDGAPGR
	1	1				GKRAAQPVAAGKVGGAAVVV
		Į.				AEPEHTKKHVKLETLLITTEGK
		l			1	LWSWGRNEKAQLGHGDTKRV
	1	l				EAPRLTEGLSHEVIVC\TCARNH
		l				TLALTETGSVFAFGENKMGQL
		l			1	GLGN*TDAVPSPVQIMYNSLPIT
		l				KMA/CGYGWLGYTEQKDEMV
						PRLVKLFDFPGHGASQIYAGYT
						CSFAVSEVGGLFFWGATNTSHE
		1			1	STMYPKAVQDLCSWRIWSLAC
					i	GKSSIIVAADESTISWGPSPTFG
		l			1	ELGYRDHKPKSSTAAQEVKTL
		l			1	HGIFSEPVAMGYSHSLVIARDE
		<u> </u>				SETEKEKIKKLPEYSPQTL

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22679	53047	A	22807		1409	INSUGLASERU WAGLGLPALPI ADEGGECGETLFVWRSQKLEG SKCKÖQLLIFGATNVDLIGRKE VPKQQAAVRNLGQNLWGPHR YGCLAGVRWTTVYSGSCAAHS LLITTEGKLWSWGRNEKGQLG HGDTKRVEAPRLIEGLSHEVIVS AACGRNHTLALTETGSVFAFGE NKMGQLGILGNQTDAVFSPAQI MYNGQPITKMACGAEFSMIMG LQRKPLFLWGALNMGQLGTLQH RWAGKPIFKMACGAEFSMIMG LQRKPLFLWGALNMGQLGTCLYPR RVAIFIEKTKDGQILPVPNVVR DVACGANHTLVLDSQKVFSW GFGGYGRIGHAEQKDEMVPRL VKLFDFFGRGASQLSCWLHTC SLCQCYNGWSVFLGGHQHLPI YHVPQKQCRTSAAGESGTLAC GKSSILVAADESTISWGPSPTFG GLGYGDHRWSSTAAGEVKTL DGIFSEQVAMGYSHSLVIARDE SETEKEKIKKLEPYNPRL
22680	53048	С	22808	277	518	
22681	53049	Α	22809	2	192	
22683	53050	A	22810	258	2946	MAEAGLRGWLLWALLIRLAQ SEPYTTIHQPGYCAFYDECGKN PELSGSLMTLSNVSCLSNTPAR KITGDHLILLQKICPRLYTGPNT QACCSAKQLVSLEASLSITKAL LTRCPACSDNFVNLHCHNTCSP NQSLFINVTRVAQLGAGQLPAV AVAEAFYQHSFAEQSVDSCSRV RVPAAATLAVGTMCGVYGSAL CNAQRWLNFQGDTGNGLAPLD ITFHLLEPGQAYGSGIQPLNEGV ARCNESQGDDVATCS
22683	53051	A	22811	258	362 4038	MAEAGLRGWLLWALLLRLAO
22004	33032		2.2012	1		SEPYTTHOPGYCAFYDEGGKN PELSGSLMTLSNVSCLSNTPAR KITGDHILLICKNICPRLYTGPNT QACCSAKQLVSLEASLSITKAL LTRCPACSDNFVNLHCHNTCSP NQSLFINVTRVAQLGAQQLPAV VAYEAFYQHSFARQSVDSCSRV RVPAAATLAVGTMGGVYGSRA CNAQRWLNPQGDTGNGLAPLD ITFHILEPGQAVGSGIQPLNEGV ARCNESQGDDVATCS

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		l		sequence		
22685	53053	A	22813	198	380	ASSESDSTPCR*MAYSWPPNSN
		l				SSHCIKELPLFMGKIPAAS*DGG
		1				PQGLAIFPPGVIVGI
22686	53054	Α	22814	1	154	
22687	53055	Α	22815	35	486	PARRGRDEC*VLAPFRATRPPS
		l				RRTR*SKATYPTPSNDQTRTIRT
		l				SRERASMPRSAEPKRPTIPRLHP
						SPRGIVTVSS/RPRSHALPVSHIH
						SCHSRTAISDNQVPAFLQVYGE
						EVRKRAGEVTQAITGFPRYRLF
						FKMLRCHRSIQVTAST
22688	53056	Α	22816	212	365	RCRSGPAPRSAPGARPASPSPPA
		1				AGTCR*RPAGHCPAGAAGLLSL
		1				CPRCST
22689	53057	Α	22817	293	618	QQNPTNCLPPASR*SKT*RARST
						AGPTRPHLHHPAAPAPGSPACA
1		l				GEGRLQR/HSPNPI.YFNWKKQR
1		1				FAAKPHFRRPQSSLNPQQWLLA
						GLLQEESSSLGTDTAPLHSLG
22690	53058	Α	22818	3	1129	FCLDDLSSAINEHLSARGRRRRS
						IVKVSLHPAVIGLKSKFLKKPD
i						QLRKLF\IGGLSFETTDESLRSHF
		1				EQ\WGTLTDCVVMRDPNTKRS
		1				RG\FGFVTYATVEEVDAAMNA
		1				RPHKVD\GRVV\EPKRAVSREDS
		1				QRPGA\HLTV\KKIFVGGIKEDT
		1				EEHHLRDYFEEIILNSMGKLEV
		1				VEIH*LDR\GQWPRKRGFAFVT
		ı				FDGHDSVDKIVIQKYHTVNG\H
1		ı				NCEVRKALSKQEMASASS\SQR
		ı			İ	G\RVGSGNFWCGRT\NGFGGND
		1			1	NFG\RGGNFSWSVVAFGG\SRG
		1				GVWIWVAVGDGYNGFG\NDGS
1		ı				HFG\GGGSY\NDFGDFH\NQSSK
1		1				FWTP*RGGNLGGQKLWPPMGG
1		ı				GG\QYFAKPTKPKVAYGRFPAA
		L				ASSYGKWPEDFN
22691	53059	Α	22819	[1	568	MGKMSPGHVRDFHGSCHHNRP
l		ı				GGPGGKSGFMGQALGPG/GCV
		1	1	1	l	QPRDLVLCVSATPAMAERGQC
			1			GAQAVASEGGSPRPWQLPCGV
						EPADFRRYMETPGCPGRNL/PA
		1		1		GAGPL*RTSAGAVWKGNVRSE
			1			SP/PQSLYWGTASGAVRRGPPSS
						RNQNGRSTDSLHCMPGKDTQ\P
		_				NP*KQPGGRLHPAKPQGRSCP
22692	53060	Α	22820	232	389	

22693 53061 A 22821 3	SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
22693 33061 A 22821 3 401 VGVCEKRGAPGIPPLAPG PVAAAVPPGPGAARAPM AAAGGRGPPEQ*T**KAW GTGSLJRGHPAPM**PLLR. PLHPLPPPHEPPRPRAAISN KCRTAAGTRQWGGRQSC QCLR	NO:						
22693 33061 A 22821 3 401 VGVCEKRGAPGIPPLAPG		sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
PVAAAVPP/GPGASRARP AAAGGRPEQ*T*FAXW GTGSL/RGHPAP*PLLR. PLHPLPPPHPLPRAAJSN KCRTAAGTRQWGGRQSC QCLR QCLR GCL					sequence		
PVAAAVPP/GPGASRARP AAAGGRPEQ*T*FAXW GTGSL/RGHPAP*PLLR. PLHPLPPPHPLPRAAJSN KCRTAAGTRQWGGRQSC QCLR QCLR GCL	00.000		 	Inner	10	I do 1	LIGUIGEN DO LEGISTRY LEGISLES
AAAGGRPEQ*T*FAXW	22693	53061	A	22821	3	401	
GTGSL/RGHPAPM*PLLR.			ı				
PLHPLPPPHPLERRAALSN KCRTAAGTRQWGGRQSC QCLR							
CRTAAGTRQWGGRQSC CQLR							
QCLR			ı				PLHPLPPPHPLPRRAA\SNPGTS
22694 53062 A 22822 1 410 MGKMSPCHVRDFHGRES RGPGGKSGYGKVQGPH LGTWCLVSQPLHFIG*KG SGVASEGGSPKPWQLPR MGAQKSRTEVKPEPLPRE GNTWMPRIPMCCARDPG GCSNPSQRR GRPHADPANYSARNLSRE GRVQDPRQAPQLPPGLL VRRPPRRPVDRTLPGGP AAAGPTGGRGSPRAGG TGPRAGFQAKGAAGSARFUN NMGRPPPRATTIPPGAA* REEAPGKRYSPPGPSGA TGPRAGFQAKGAAGSARFUN TGRAGFGAKGAGGGDS ILLVADAGPDSARSPPGL LSAAPRSRRDPDAVGGL LSAAPRSRRDPDAVGGL LSAAPRSRRDPDAVGGL LSAAPRSRRDPDAVGGL LSAAPRSRRDPDAVGGL LSAAPRSRRDPDAVGGL AARRINTTWKLPQLGAS CKMVDCLANSEANTRRISI FGAAGQPLTIPGRVLIGGE KLVCKKPKARQFFRIEND GNIVIGKKYKIQHIPLE SIKDEGDLRNGWVIKTPT VYAAATATEKSEWMNIN DLLSKSGKTPSNEHAAVV SEATVCMRCQKAKFTPV HCRKGGFVVGPCSEKRI QSSKPVRICDFCYDLLSA TCQPARSDSYSQSLKSPL DDDDDDDMQ DDDDDDMQ DDDDDDMQ DDDDDDMQ DDDDDDMQ DDDDDDMQ DDDDDDMQ DRTNIMVGYHQTNQKTD PTIRRPVLYDHNRLPEGSS AVAVAGDHPAAVQVTM DTGFPPVFSGSIAESWGG PSY*CWBATMILRAFPI GETKDLRQFPNIDNAYMI							KCRTAAGTRQWGGRQSQLQRC
RGPGGKSGFVKVQGPH LGTWCLVSQFLHPiQ*KG SGVASEGGSFKPWQLPR MGAQKSRTEVWEPLPRE MGAQKSRTEVWEPLPRE GNTWMPRIPMCARDEG GCSNPSQRR GCSNPSQRR GPRHADPANFSARNLSRT GRVQPPRPQAPQLIPPGLVRPPPAPQAPQLIPPGLVRPPPRPVDRTLPGGP AAAAOPTGGRSGSPRAGG TGPRAGPARQAAAOPTGGRSGSPRAGG TGPRAGPARQAAQAAAOPTGGRSGSPRAGG TGPRAGPARQAAQAAAAOPTGGRSGSPRAGG TGPRAGPASRAGAGSAAGSAP NMGRPPPATTIPPGAA* REEAPGKFRVSPPGPSGA TRASFITIAPGGAGGGDSB ILLVADAQPDSARSPPGL LSAAPRSRRDPPDAVGGL AARRINITWKLPQLGAS LSAAPRSRRDPPAVGGL AARRINITWKLPQLGAS GRIVQKKYKIKQHIPLE SIKDEGDLRNGWVIKTPT VYAATATEKSEWMNHIN DLLSKSGKTPSNEHAAV SEATVCMRCQKAKFTEV HCRKGGFVCPSEKRI QSSKPVRICDFCYDLLSA TCOPARSDSVSQLKSPLJ DDDDDDDMQ DDDDDDMQ DDDDDDMQ DDDDDDMQ DRTNIMVGYHQTNQKTD PTRRPVLVDHNRLPEGSS AVAVAGDIPAAVQVTM DTGFDPVFSGSIAESWGQ PSYC*DWRATMLRAPF GETKDLROFFNIDNAYMI GETCHLORY GETCHL							
LGTWCLVSQPLHPIG*KG SG\VASEGSPKPWQLPR MGAQKSRTEVWEPLPRF GNTWMPRIPMCCARDPG GCSNPSQRR GCSNPSQRR GCSNPSQRR GPRHADPANFSARNLSR GRVQDPRQAPQLPPCL VRRPPRRPVDKTLPGGP AAAGPTGGRGSPRAGG TGPRAGPGARGAGASAP NMGRPPPATTTPPGAA* REEAPGKRVSPPPGASAGAGAP NMGRPPPATTTPPGAA* REEAPGKRVSPPPGASAGAGAP NMGRPPPATTTPPGAA* REEAPGKRVSPPPGASAGAGAP NMGRPPPATTTPPGAA* REEAPGKRVSPPPGASAGAGAP NMGRPPPATTTPPGAA* REEAPGKRVSPPPGASAGAGASAP NMGRPPPATTTPPGAA* REEAPGKRVSPPPGASAGAGASAP NMGRPPATTTPPGAA* REEAPGKRVSPPPGASAGAGASAP NMGRPPATTTPPGAA* REEAPGKRVSPPGASAGAGASAP NMGRPPATTTPPGAA* REEAPGKRVSPPGASAGAGASAP NMGRPPATTTPPGAA* REEAPGKRVSPPGASAGAGAGASAP NMGRPPATTTPPGAA* REEAPGKRVSPPGASAGAGAGAGABAGAGAGAGAGAGAGAGAGAGAGAGAGA	22694	53062	Α	22822	1	410	MGKMSPGHVRDFHGRPSHHRP
SGVASEGGSFKPWQLPR MGAQKSRTEVWEPLPRF GNTWMPRIPMCCARDFG GCSNPSQRR GCSNPSQRR GCSNPSQRR GGSNPSQRR GFRHADPANFSARNLSRF GRVQDPRQAPQLPPGL VRRPPPRRPVDRTLPGGP AAAACPTGGRGSPRAGG TGPRAGPGARGAAGSAP NMGRPPPATTTPPGAA* REEAPGKRVSPPGPSGA TRASPTPTAAGGGGOSS II.LVADAGPDSARSPPGL LSAAPRSRRDPDA VGGLA AARRINTTWKLPQLGAS II.LVADAGPDSARSPPGL LSAAPRSRRDPDA VGGLA AARRINTTWKLPQLGAS GKMVDCLANSEANTRRISI FGAAGQPLTIPGRVLIGEG KLVCRKKPKARGFELRRD GRIVQKKKYIKQHIPLE SIKDEGDLRGWVIKTPT VYAAATATEKSERWMHIN DLLSKSGKTPSNEHAAVV SEATVCMRCQKAKFTEV HCRKGGFVVCPSEKR QSSKPVRICDFCYDLLSA TCOPARSDSYSQLKSPLJ DDDDDDDMQ DDDDDDMQ DDDDDDDMQ DDDDDDMQ DRTNIMVGYHQTNQKTD PTRRPVLYDHNRLPEGSS AVAVAGDIPBAAVQVTM DTGFDPVFSGSIAESWGQ PSY***DWFAATMLRAFP GETKDLRGFRUDNAYMI GETKDLRGFRUDNAYMI DGFDPVFSGSIAESWGQ PSY****DWFAATMLRAFP GETKDLRGFRUDNAYMI GETKDLRGFRU							RGPGGKSGFVGKVQGPHAVCS
MGAOKSRTEVWEPI.PRE							LGTWCLVSQPLHP\G*KGPMYS
GNTWMPRIPMCCARDPG			1				SG\VASEGGSPKPWQLPRGVEP
GCSNPSQRR							MGAQKSRTEVWEPLPRFQKIY
22695							GNTWMPRIPMCCARDPGGAKG
GRVOPPRQAPQLIPPGL VRRPPPRRPVDRTLEGE AAA/OPTGGGSSPRAGG AAA/OPTGGGSSPRAGG AAA/OPTGGGSSPRAGG TGPRAGPGARGAAGSAP NMGRPPPATTIPPGAA* REEAPGKRVSPPGPSGA TRASPIPTAAGGGGGDS IILVADAG/PDSARSPPGL LSAAPRSRCPPDAVGGL/ AARRINITWKLPQLGAS LSAAPRSRCPPDAVGGL/ AARRINITWKLPQLGAS EVALUATION AARRINITWKLPQLGAS FGAAGQPLTIPGRVLIGE KLVCKKPKARGPELRID GRIVINGKKYKIKGHIPLE SIKDEGDLRIGWVIKTPT VYAATATEKSEWMHIN DLLSKSGKTPSNEHAAV SEATVCMRCQKAKFTEV HCRKGGFVCPGSEKR QSSKPVRICDFCYDLLSA TCQPARSDSYSQSLKSPLI DDDDDDDMQ 22698 \$3066 A 22826 1219 2370 22698 33067 A 22827 164 601 DRTNIMVGYHQTNQKTD PTIRRPULDHNRLPEGSS AVAAGDIPAAVQVTM DTGFDPVFSGSIAESWGQ PSYC*DWEATMLRAFP GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GROFFNOM GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GROFFNOM GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GETKDLROFFNDINAYMING GROFFNOM G							GCSNPSQRR
VRRPPPRRPVDRTLPGGG AAA(GPTGGRGSPRAGG AAA(GPTGGRGSPRAGG TGPRAGCARGAAGAAGAP NMGRPPPATTTPFCGAA- REEAPGKFRVSPPGPSGA TRASPTPTAAGGGGGDS II.LVADAGPDSARSPPGL LSAAPRSRRDPDAVGGL/ AARRINTTWKLPQLGAS A 22824 271 426 KMVDCLANSEANTRRISI FGAAGQPLTIPGRVLIGEC KLVCRKKPKARGFERND GNIVIQKKVIKQHIPLE! SIKDEGDLRNGWVIKTP VYAATATEKSEWMHIN DLLSKSGKTPSNEHAAVV SEATVCMRCQKAKFTEV HCRKCGFVVCGPCSEKRI QSSKFVRICDFCYDLLSA TCQPARSDSYSQSLKSPLI DDDDDDDMQ 22699 53067 A 22826 1219 2370 DRTNIMVGYHQTNQKTD PTRRPVLOPINRLPEGS AVAYAGDIFBAAVQVTM DTGFPPVFGSIAESWGG PSYC*DWEAATMLRAFPI GETKDLRQFFNIDNAYMI GROUNT G	22695	53063	A	22823	264	979	GPRHADPANFSARNLSRP*DAP
VRRPPPRRPVDRTLPGGG AAA(GPTGGRGSPRAGG AAA(GPTGGRGSPRAGG TGPRAGCARGAAGAAGAP NMGRPPPATTTPFCGAA- REEAPGKFRVSPPGPSGA TRASPTPTAAGGGGGDS II.LVADAGPDSARSPPGL LSAAPRSRRDPDAVGGL/ AARRINTTWKLPQLGAS A 22824 271 426 KMVDCLANSEANTRRISI FGAAGQPLTIPGRVLIGEC KLVCRKKPKARGFERND GNIVIQKKVIKQHIPLE! SIKDEGDLRNGWVIKTP VYAATATEKSEWMHIN DLLSKSGKTPSNEHAAVV SEATVCMRCQKAKFTEV HCRKCGFVVCGPCSEKRI QSSKFVRICDFCYDLLSA TCQPARSDSYSQSLKSPLI DDDDDDDMQ 22699 53067 A 22826 1219 2370 DRTNIMVGYHQTNQKTD PTRRPVLOPINRLPEGS AVAYAGDIFBAAVQVTM DTGFPPVFGSIAESWGG PSYC*DWEAATMLRAFPI GETKDLRQFFNIDNAYMI GROUNT G			l				GRVQDPRPQAPQL\PPGLGAPSR
AAA/ADPTGGRGSPRAGG: TGPRAGPGARGAAGSAP NMGRPPPATTITPPGAA* REEAPGKRYSPPGPSGA TRASPTPTAGGGGGDS ILLVADAGPDSARSPPGI LSAAPRRRDPDA VGGL AARRINTTWKLPQLGAS A 22824 271 426 Z2697 \$3065 A 22825 \$46 1296 KMVDCLANSEANTRRISI FGAAGQPLTIPGRVLIGGE KLVCKKPKARQFFLRND GNIVIGKKYIKQHIPLE! SIKDEGDLRNGWVIKTPT VYAATATEKSEWMNIN DLLSKSGKTPSNEHAAVV SEATVCMRQKAKFTPV HCRKGGFVVGPCSEKRI QSSKPVRICDFCYDLLSA TCQPARSDSYSQLKSPLJ DDDDDDDMQ DDDDDDMQ Z2699 \$3067 A 22826 1219 2370 DRTNIMVGYHQTNQKTD PTRRPVLYDHNRLPEGS AVAVAGDHPAAVQVTM DTGFPPVFSGSIAESWG GPKVBCATMLRAFPI GETKDLRGFPNUDNAYMI GETKDLRGFPNUDNAYMI GETKDLRGFPNUDNAYMI GETKDLRGFPNUDNAYMI GETKDLRGFPNUDNAYMI GETKDLRGFPNUDNAYMI GETKDLRGFPNUDNAYMI GETKDLRGFPNUDNAYMI GETKDLRGFPNUDNAYMI GETKDLRGFPNUDNAYMI TGAAPTT TRASPTCA TASPTT TASPT			ı			Ì	VRRPPPRRPVDRTLPGGPRPPA
TGPRAGPCARGA/GSAP NMGRPPPPATTIPPGAA* REEAPGKFRVSPPGPSGA TRASPIPTAAGGGGGDS II.LUVADAGVPSARSPPGI LSAAPRSRDPPDA/GGL/ AARRINTTWKLPQLGAS LSAAPRSRDPPDA/GGL/ AARRINTTWKLPQLGAS S3065							AAA\GPTGGRGS\PRAGGTAAA
NMGRPPPATTTPPGAA REEAPGKFRVSPPPGAA REEAPGKFRVSPPPGAA REEAPGKFRVSPPPGAA TRASFITIAAGGGGDS II.LVADAGPDSARSPPGL LSAARSRRDPDA VGGLL AARRINTTWKLPQLGAS A 22824 271 426 ARRINTTWKLPQLGAS A 22825 546 1296 KMVDCLANSEANTRRISI FGAAGQPLTIPGRVLIGE KLVCRKPKARGFERND GNIVIQKKKYIKQHIPLE SIKDEGDLRRGWVIKTPT VYAATATEKSEKWMNIN DLLSKSGKTPSNEHAAVV SEATVCMRCQKA KFTEV HRKGGFVVCGPCSEKRI QSSKPVRICDFCYDLLSAA TCQPARSDSYSQLKSPLI DDDDDDDMQ DDDDDDMQ 22699 53067 A 22827 164 601 DRTNIMVGYHQTNQKTD PTRRPVLVDHNRLPEGSF AVAVAGDIFBAAVQVTM DTGFPPVFSGSIAESWGQ PSYC*DWEAATMLRAFPI GETKDLRQFPNIDNAYMI							
REEAPGKERVSPPGPSGA			1				
TRASPIPIAAGGGGGDS IILVADAGPDARSPPRI LESA PRISREPPO LESA PRISREPPO LESA PRISREPPO LESA PRISREPPO LESA PRISREPPO LESA PRISREPPO LESA PRISREPPO LESA PRISREPPO LESA PRISREPPO LESA PRISREPPO LESA PRISREPPO LESA PRISREPPO LESA PRISREPPO LESA PRISREPPO LESA PRISREPI LESA P			1				
II.LVADAGIPDRARSPPGL			1				
LSA APRSRROPDA VGGLA			Į.]		
AARRINTTWKLPQLGAS 22826 271 426 KMVDCLANSEANTRRISI FGAAGQPLTIPGRVLIGEG KLVCRKPKARAGPERNID GNIVIQKKYIKQHIPLE SIKDEGDLRNGWVIKTPT VYAATATEKSEWMYHIN DLLSKSGKTPSNEHAAVV SEATVCMRCQCAK FTPV HCRKCGFVVCGPCSEKRI QSSKPVRICDFCYDLLSA TCQPARSDSYSQLKSPLI DDDDDDDMQ 22699 53067 A 22827 164 601 DRTNIMVGYHQTNQKTD PTRRPVLYDHNNRLPEGSF AVAVAGDIFBAAVQVTM DTGFPPVPSGSIAESWOG PSYC*DWEAATMLRAFPI GETKDLRQFPNIDNAYMI			1		ļ	İ	
22696 \$3064 A 22824 271 426	1	j	1				
22697	22606	52064	ļ.	22824	271	426	AARRENTTWKEPQEGASTE
FGAAGQPLTIPGRVLIGEC KLVRKKPKARGFTLAGE KLVRKKPKARGFTLAGE KLVRKKPKARGFTLAGE KLVRKKPKARGFTLAGE SIKDEGDLRNGWVIKTPT							MANDOLANGEANT DRIGIVENO
KLICRKKPKARQFFLRND GNIVIQKKKYIKQHIPLE SIKDEGDLRNGWVIKTPT	22097	33063	^	22023	346	1290	
GNIVIOKKY/IKQHIPILE: SIKDEGDLRNGWVIKTPT VYAATATEKSEWMNHIN DLLSKSGKTPSNEHAAVV SEATVCMRCQCAKFTPV- HCRKCGFVVCGPCSEKRI QSSKPVRICDFCYDLL5A- TCOPARSDSYSQLKSPLI DDDDDDDDMQ 22698 53066 A 22826 1219 2370 22698 53067 A 22827 164 601 DRTNIMVGYHQTNQKTD PTRRPVLOMNRLPEGSE AVAYAGDHFAAVQVTM DTGFPPVFSGSLAESWGQ PSYC*DWEAATMLRAFPI GETKDLROFFNIDNAYMI		i	1				
SIKDEĞDLRAĞW'İKTPT VYAATATEKSEWMNHIN DLLSKSĞKTESNEHAAV\ SEATVCMRCQKAKFTEV\ HÇRKĞĞFVVÇPĞSEKRİ QSSKPVRICDFCYDLLSA TCOPARSDSVSQSLKSPLI DDDDDDMQ		ł	1				
VYAATATEKSEWMNIND			1				
DLLSKSGKTPSNEHAAVV SEATVCMRQKAKFTPV HCRKGGFVVGPCSEKRI			1				
SEATVCMRCQKAKFTPV HCRKCGFVVCGPCSEKRI			1				
HCRKGGFVVČQPCSEKRI			1				
QSSKPVRICDFCYDLLSAG TCQPARSDSYSQSLKSPLI DDDDDDDMQ 22699 53067 A 22826 1219 2370 DRTNIMVGYHQTNQKTD PTRRPVLVDHNRLPEGSF AVAVAGDHPAAVQVTM DTGFPPVFSGSIAESWGQ PSYC*DWEAATMLRAFPI GETKDLRQFPNIDNAYMI			1				
TCQPARSDSYSQSLKSPL1		l	1				
DDDDDDDMQ		l	1				QSSKPVRICDFCYDLLSAGDMA
22698 53066 A 22826 1219 2370 DRTNIMVGYHQTNQKTD			1				TCQPARSDSYSQSLKSPLNDMS
22699 53067 A 22827 164 601 DRTNIMVGYHQTNQKTD PTRRPULVOHNRLPEGSI AVAYAGDIFBAAVQVTM DTGFDPVFSGSIAESWGQ PSYC*DWEAATMLRAFPI GETKDLRQFPNIDNAYMI						l	DDDDDDDMQ
PTRRPVLVDHNRLPEGSF AVAVAGDHPAAVQVTM DTGFDPVFSGSIAESWGQ PSYC*DWEAATMLRAFPI GETKDLRQFPNIDNAYM							
AVAVAGDHPAAVQVTM DTGFPPVFSGSIAESWOG PSVC*DWEAATMLRAFPI GETKDLRQFPNIDNAYMI	22699	53067	A	22827	164	601	DRTNIMVGYHQTNQKTDTGKT
DTGFPPVFSGSIAESWGG PSYC*DWEAATM.RAFPI GETKDLRQFPNIDNAYMI			1				PTRRPVLVDH\NRLPEGSRGRL
PSYC*DWEAATMLRAFPI GETKDLRQFPNIDNAYMI	1		1		1	I	AVAVAGDHPAAVQVTMTLVN
GETKDLRQFPNIDNAYME			1				DTGFDPVFSGSIAESWGQHSGT
		l	1		1	1	PSYC*DWEAATMLRAFPLAKK
NRADAVI HDTPNII YEME	1	l	1				GETKDLRQFPNIDNAYMELGT
I I I I I I I I I I I I I I I I I I I		l	1				NRADAVLHDTPNILYFMQF
22700 53068 A 22828 2 91	22700	53068	Α	22828	2	91	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
i	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22701	53069	Α	22829	1	1188	MGAEPSLLTSLGTLWSGERFOC
		i				MRAKIWERSRKGRRFSTDVISP
1						TKDSFAGPFQPISDTLVQHQRIY
ŀ	İ	ĺ				GLKVCVLPDPCKIRILKPNPQC
İ		ŀ				VGKWEEEVAENEIQVFGGERF
	İ		l			DINKYSKSPPPPTPQPLYLTAGF
ŀ		ı				ASCRNHRKSEELREGKGESPGK
		l				SKPFHLQANFTVALVTTLQGDK
						HSQEAFIRPVAKAKTTCTYLSN
						VCEMKDFKRINCPDPFHKMAP
	1	1				KAKKEA\PAPP\KAEAKAKGFK
1						GPRRQVLK\GVHNHKKRKRKK
		1				ERSRTFTHLSRRPGRPLRI\RRQP
	l	1				KYPSGRAAPRRNKLD\HYAIIK
		i	1			FPLT\TESA\MKK1EDNNTL\VFI\
		l				VELLKATTKNQIKQGC*RSLYD
1		1				I\DVGQGSTPWISAWMGEKKGI
						CSQLAPDLPIAFGMFANKIWGFI
22702	53070	A	22830	1413	2197	RVEWRWYNVLPCLDVHSGPVD
1						RRSHVMTAAEKPPFCPAWGQA
İ		1				VSQQSCALEPRTGEEGGKGIQG
			ŀ			PASGLFPTGEEISVVPNRWLDA
1						SLSPVILGVQG\GSQCLSCGVGQ
ŀ						EPTLTLEVRLGASSLGTQPQML SLLKPGSPQPWCCGTP*QDSVD
ľ		l				GSFASSLRILAKALLCGPGFCPS
l						LTLPSFCRKPVTIME\LYLGAKE
ŀ		l				S\KSFTFYRRDMG\LT\SSFESAA\
ŀ		l				YPGWFLCTEPE\ADOPVRLTOL
	ŀ	l				PENGGWNAPITDFYFQQ\CD
22703	53071	A	22831	100	177	KRDHF*SRLQQVSRSVRWSSSA
22/03	33071	ľ	22031	100	l'''	TDSR
22704	53072	A	22832	610	1080	1DSK
22704	53072	A	22833	1169	1308	
22703	22013	<u></u>	22033	1107	1300	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop eodon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequenee	deletion, \=possible nucleotide insertion)
				sequenec		
22706	53074	Α	22834	l	1889	ACRRRAGTWTPPRPRGSGSRA
						MAGNCGARGALSAHTLLFDLP
		l				PALLGELCAVLDSCDGALGWR
			1			GLAERLSSSWLDVRHIEKYVDQ
						GKSGTRELLWSWAQKNKTIGD
	İ					LLQVLQEMGHRRAIHLITNYGA
	l	ĺ				VLSPSEKSYQEGGFPNILFKETA
						NVTVDNVLIPEHNEKGVLLKSS
						ISFQNIIE\GTRNFHKDFLIGERG
		i				E/IFGGYQEGKIQTLTYAVKLFK
						QEKKMQCKKHWKRFLSELEVL
						LLFHHPNILELAAYFTET\EKFC
					,	LIYPYMRNGTLFDRLQCVGDT
						APLPWHIRIGILIGISKTIHYLHN
						VQPCSVICGSISSANILL\DDQFH
						PKLTDFSMAHFRSP\LEHQSCCI
						NMPSS\SRKHLWY\MPEEYIRQ
						GKLSIKTDVYSFGIVIMEVLTGC
						RVVLDDPKHIQLRDLLRELMEK
						RGLDSCLSFLDKKVPPCPRNFS
						AKLFCLAGRCAATRAKLRPSM
						DE\V*RALESTQASLYF\AGDPP
						TSLKALRGP\SPLF\LEKVT\SIPV
						\EDDE\KPNNNLLPS**KALRMK
						RWPQK\TPF\ECSQP\EVWFLSLE
						QKAREARENEEACNMPSSS\CE
		l				ESWFPKY\IVPS\QDLRPL*GK*K
						ILSSQA\SGHSC\RTKPVESSCSS
						KFSWDEYEQYKKE

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Uaknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22707	53075	A	22835	1	1560	RWERGGKRFRAADPMSALRRS GYGPSDGPSYGRYYGPGGGDV PVHPPPPLYPLRPEPPOPPISWR
1		1				VRGGPAETTWLGEGGGGDG
		l				YYPSGGAWPEPGRAGGSHOEO
	1	ı				PPYPSYNSNYWNSTARSRAPYP
		l				STYPVRPE\LQGQSLNSYTNGA
	1	l				YGPTYPPGPGANTASYSGAYY
		1		1		APGYTQTSYSTEVPSTYRSSGN
1	i	l				SPTPVSRGIYPOODCOTEAPPLR
1	1	1				GQVPGYPPSQNPGMTLPHYPY
		l				GDGNRSVPQSGPTVRPQEDAW
		1				ASPGAYGMGGRYPWPSSAPSA
	1	ı				PPGNLYMTESTSPWPSSGSPQSF
		l				PSPPVQQPKDSSYPYSQSDQSM
		l				NRHNFP/CASVHOYESLGDO*T
		l				MDDSRILLDSPSPSIVPEPSACM
	İ	ı				GNAHPVTHP\NNODOSSSLPEE
		1				CVPSDESTPPSIKKIIHVLEKVO
		l	1			YLEQEVEEFVGKKTDKAYWLL
		1	1			EEMLTKELLELDSVETGGQDSV
1		l				RQARKEAVCKIQAILEKLEKKR
1		1				IMKGFITK WKPVTNLTKEHLIW
		ı		1		LITLFLKCLLMTRSNTFQLFL
22708	53076	A	22836	8	207	TPVPRALRSPAGLEGSHLSSIST
		1				MMPAVMSVGKVTENGGSPQGI
						KSPSKPPGPNRIG*RNQVSVHT
22709	53077	Α	22837	459	5520	RQVRPAGLEGSDLSSINTMMSA
	1	ı				VMSVGKVTENGGSPQGIKSPSK
1		l		l]	PPGPNRIGRRNQETKEEKSSYN
i				ŀ		CPLCEKICTTQHQLTMHIRQHN
		l				TDTGGADHSCSICGKSLSSASSL
		l				DRHMLVHSGERPYKCTVCGQS
		l				FTTNGNMHRHMKIHEKDPTSA
		1				TATAPGRGDFASAKSSKRKLSH
	1	1				DAESEREDPAPAKKMVEDGQS
		1				GDLEKKADEVFHCPVCFKEFV
		1				CKYGLETHMETHSDNPLRCDIC
22710	62070	١.	22838	164	424	CVTFRTHRGLLRHNAL
22710	53078	A	22839	164	242	
22711	53080	A	22840	187	325	
22713	53081	A	22841	1	130	NCPMCWPGOSPRVPST*GRRNP
[227.13	23001	ľ	22041	ľ	1.50	PPKRRSVSAES*RHWPQSPSP
22714	53082	A	22842	614	774	OEPMRMCTMLANSSPSSSOATP
-2//		Γ.		1	l'''	PGL*SWNQPSLPARTLSRHEWT
			l		1	SCLGSSANL
22715	53083	A	22843	396	560	LFPHDGIKHKGKENP/PKGPPPP
	1	1		I		THPOTPPRPADAPLVATKSTLD
						OAWALLAVATS
22716	53084	В	22844	124	205	

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop eodon, /=possible nucleotide
	sequence	1	09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22717	53085	-	22845	1	2355	MVGSRRISGVIPLPFTRRAGIFS
22/1/	23083	A	22843	1	2333	
		1				DPRANMQPLWAPVPPATFVPTS
	İ					LPPVPTARPPTSASCHCQQVPC
		i				AGLEVALPGNRTICWLSLWPS
	İ	1				WQLDEDAQSPALESSSIRQFPT
	l	1				APRSGYPGTLCPSFLFGPRPRTC
	ŀ					EYHSKPVLCAPSSRASPSSATTI
		1			1	STRTLWGPGAGSHPFGVHNTR
						LSPDLCPGKIVLRALKESGAGM
		1				PEQHKDPRVQENPDDQRTVPE
		l		i	į.	VTGDARSAFWPLRDNGGPSPFV
						PRPGPLQTDLHAQSSEIRYNHTS
					ŀ	QTSWTSSSTKRNAISSSYSSTGG
		1				LPGLKQRRGPASSRCQLTLSYS
						KTVSEDRPQAVSLGHTRCEKG
			İ			ADTAPGOTIAPTASRPHINTSLH
		1			1	VEDKAISDCRPSRPSHTLSSLAT
İ		l				GASGGPPVSKAPTMDAQQDRP
		1			ļ	KSQDCLGLVAPLASAAEVPAT
		1				APVSGKKHRPPGPLFSSSDPLPA
		1				NSSHSRDSAQVTSMIPAPFTAA
		1			1	SRDAGMRRTRSAPAAAAAAPP
		1				PSTLNPTSGSLLN\EWMEAPHIS
						WPOPOLOHVPRGO/NVRYNOR
		1				SQTSRTRS/CPQTKCQLQLPQLY
l						GRPPGSKAEEG\PASSHCQLALS
		1				SSNTVSEDGPQAVSSGHRCENK
		l				AGTAPGQTLAPRGGSPRSQASR
1		l				PHINTALHVEDKAISDCRPSRPS
1		1				
						HTLSSLATGASGGPPVSKAPTM
1						DAQQDRPKSQDSLGLLAPLASA
		ļ		l		AEVPSTAPVSGKKHRPPGPLFSS
22718	53086	Α	22846	1	1200	MPEQDKDPRVQENPDDQRRVP
				1		EVTGDARSAFRPLRDNGGLS/PF
1						FVPGPGPLQTDLHTQR\LGYRV
1		1		l		TVEDLDREKEAAFQRINSALQV
1				l	i	EDKAISDCRLSRPSHTLSSLATG
			ł	l		TSGLPAVSKAPSMDAQQERHK
l				l		SQDCLGLLAPLAPAAEVPSTAP
		l		l		MSGKKHRPPGPLFSSSDPLPATS
1						SHSQDSAQVTSLIPAPFPAASM
				l		DAGMRRTSPGTSAPAAAAAPPP
1						STLNPTLGSLLEWIEALHISGPQ
1	1	1	1	I		PQLQQVPRGQNQRSQTSRTSSC
	1	1	1	1	1	PKRK/CHLQLLQLYGRPPGTKA
						EEG\PASSHCQLTLSSSNTVSED
1	1		l	1		GPQAVSSGHTHCEKTADTAPG
1	1	1	i			QTLAPRGGFPRSQASRPCRHKF
		1	l	l		PLLPRRRGEPLMLPPPLELGYR
		1				VTAEDLDREKEVAFQRIKSALQ
1	1	1		1		VEDKAI
					l	VEDRAI

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22719	53087	Α	22847	2	1762	YHSKPVLCAPSSRASPSSATTLS
						TRTLWGPGAGSHPFGVHNTRL
		1				SPDLCPGKIVLRALKESGAGMP
ļ	i					EQHKDPRVQENPDDQRTVPEV
						TGDARSAFWPLRDNGGPSPFVP
		ŀ				RPGPLQTDLHAQSSEIRYNHTS
ł						QTSWTSSSTKRNAISSSYSSTGG
		ŀ				LPGLKQRRGPASSRCQLTLSYS
		1				KTVSEDRPQAVSLGHTRCEKG
		ŀ				ADTAPGQTIAPTASRPHINTSLH
		1				VEDKAISDCRPSRPSHTLSSLAT
l		ŀ				GASGGPPVSKAPTMDAQQDRP
		[KSQDCLGLVAPLASAAEVPAT
l		l				APVSGKKHRPPGPLFSSSDPLPA
	ł	1				NSSHSRDSAQVTSMIPAPFTAA
		1				SRDAGMRRTRSAPAAAAAAPP
		ł				PSTLNPTSGSLLN\EWMEAPHIS
ŀ	İ					WPQPQLQHVPRGQ/NVRYNQR
	1	l				SQTSRTRS/CPQTKCQLQLPQLY
ļ				1		GRPPGSKAEEG\PASSHCQLALS
	İ	1		ļ		SSNTVSEDGPQAVSSGHRCENK
	ł	1				AGTAPGQTLAPRGGSPRSQASR
ŀ		1				PHINTALHVEDKAISDCRPSRPS
				1		HTLSSLATGASGGPPVSKAPTM
			ł		}	DAQQDRPKSQDSLGLLAPLASA
					f	AEVPSTAPVSGKKHRPPGPLFSS
1						SDPLPATSYHSRDTAQ

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22720	53088	A	22848	292	2053	GSVAWSPFSRSSLCLWDLPV APTLWGPGAGSHPFGVHNTRL SPDLCPGKIVLRALKESGAGMP EQDKDLRVQENPDDQRRVPEV TGDARSAFLAPAGQWSPLSLC AQSRASADIPPCFEVRQI*PDIP HLDELMIPOKCHLQLLQLCG RLAGAKVEEGASGAKEQGRDA "AGQGTPESQQNPDAQRTVPEV RGPLQTDLHAQSSEIRYNQTS GTSWTSSTKRNAISSSYSSTGG LIGLKQRRGPASSRCQLTLSYS KTVSEDRPQAVSSGHTRCEKA ADTAPGQTLAPRGGSPRSQGSR PRRRKIALLPHR*GELQVEDKAI LDCRFSWPSHTLSSLATGASGE PPVSKAPTMDAQQDRPKSQDC LLGLVAPLASAAGIPSTAPVSGK KHRPPGELFSSSDPLPATSSHISS DSAQVTSLIPAAFTAASMDVG
						LNPTSGSLLN\EWMEALHISGPQ PQLQQVPRGQ/NVRYNQRSQTS RTRS/CPQTKCQLELPQLYGRPP
						GTKGCRHEKNVLCSKLFEGFGF IFVGFFFFAYMGILQLLII

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22721	53089	A	22849	146	1789	RQPSTSGSSRSCPCARRSWWAR
		l				TARSGWWWELLGARRSPOTLH
		l				CPQATLITLPCPRPSSTTSGSAM
		ŀ				T*RGPWRSPGCTTSFCPTSR/PV
		1				ERNIDQTDLHAQRSEIRYNQTS
		1				QISWTSSCTNRNAISSSYSSTGG
		i				LPGLKRRRGPASSHCQLTLSSS
		1		1		KTVNSDKFDANDPILKDQTQE
l		1				WSGSATFTSDGKIRLFYTDYSG
						KHYGKQSLTTAQVNVSKSDDT
ĺ						LKINGVEDHKTIFDGDGKTYQN
İ						VQQFIDEGNYTSGDNHTLRDPH
		l				YVEDKGHKYLVFEANTGTENG
						YQGEESLFNKAYYGGGTNFFR
		1				KESQKLQQSAKKRDAELANGA
1		1				LGIIELNNDYTLKKVMKPLITR
		ı				GSSRPSIREDENGRSQKPVHPEG
						DMTMNIKKIVNQATVLTFTTAL
		1				LAGGATQAFAKENNQKAYKET
		1				YGVSHITRHDMLQIPKQQQNEK
		1				YQVPQFDQSTIKNIESAKGLDV
		l				WDSWPLQNADGTVAEYNGYH
		1		1		VVFALAGSPKDADDTSIYMFY
		İ			1	QKVGDNSIDSWKNAGRVFKDS
		L				DKFDANDPILKDQTQEWRGEA
22722	53090	A	22850	500	1034	

SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO: in USSN	Nucleotide Jocation of first	Nucleotide location of last	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO.	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22723	53091	Α	22851	1	2396	MDKYLLHGPPRPPPAIASRCLA
						RVPGCAYSSIMVALSLVPLCAH
		1				MPTPSGLEVALPGNRTICWLSL
		1				WPSWQLDEDAQSPALESSSIRQ
		l				FPTAPRSGISPLCAPHSCLGQGF
	Ì	l				GPASHPTSTQTLDISGVKTTPRI
		l				FSVILQDLCSILGDSCLQFHLHR
	ļ.	l	Į.			VPQQVSAVLLRVSGASPSPATT
						LSTRTLWGPGAGSHPFGVHNT
	ł	1				RLSPDLCPGKIVLRALKESRAG
		l				MPEQDKDPRVQENPDDQRTVF
						EVTGDARSAFWPLRDNGGPSP.
	1		İ			VPRPGPLQTDLHAQSSEIRYNH
						TSQTSWTSSNTKRNAISSSYSST
						GGLPGLKQRRGPASSRCQLTLS
	l	1				YSKTVSEDRPQAVSSGHTRCE
		i				GADTSPGQTIAPTGGSPRSQDS
						RPRRKIPLLPRRRGEPLMLPPF
	ļ	l		1		LELGYRVTAEDLHLEKETAFQI
				1		INSALHVEDKAIPDCRPSQPSH
	l					LSSLATGASGGPPVSKAHTMD
						AQQHRPKSQDCLGLLAPLASA
						AEVPSTAPVSGKKHRPPGPLFS
		1				SDPLPATSYHSRDSAQVTSLIPA
		l		l		PFTAASSDAGMRRTRPGTSAPA
	l			ŀ		AAAAAPPPSTLNPTSGSLLN\EV
						MEAPHIS WPQPQLQHV SRGQ/N
				1		VRYNQRSQTSRTRS/CPPRKCQ
				1		QLPQLYGRPPATKAEEG\PASSI
	l .			1		CQLAHSSSNTVSEDGPQAVSSC
				1		HTRCEKKAGAS/RWASRF*STH
		_				YGCTAGQTQVPRLPGPTGPPSI
22724	53092	A	22852	1	3392	MDKYLLHDPPRPPPAIASRCLA
	i					RVPGCAYSSIMVALSLVPLCAF
		1			1	MPTPSGLEVALPGNRTICWLSL
		1				WPSWQLDEDAQSPALESSSIRC
		ı				FPTAPRSGYPGTLCPSFLFGPRP
						RTCEYHSKPVLCAPSSRASPSS/
		l				TTLSTRTLWGPGAGSHPFGVHI
		1				TRLSPDLCPGKIVLRALKESGA
		1				GMPEQDKDPRVQENPDDQRT\
		1				PEVTGDARSAFWPLRDNGGPSI
			l			FVPRPGPLQTDLHAQSSEIRYN HTSOTSWTSSST
	1	1	l			
22726	52002	_	22062	40	400	
22725	53093	A	22853	40	409	WKMRHETRYNVLVSRGVGCH
22725	53093	A	22853	40	409	WKMRHETRYNVLVSRGVGCH RLFFLNGVFEHLGRAFVQRRK
22725	53093	A	22853	40	409	WKMRHETRYN VLVSRGVGCH RLFFLNGVFEHLGRAFVQRRKV DTEHQLLRAA VPANQEHRGLT
22725	53093	A	22853	40	409	WKMRHETRYN VLVSRGVGCH RLFFLNGVFEHLGRAF VQRRK V DTEHQLLRAA VPANQEHRGLT QSQHVDGHSLLLLTLPNSFLCQ
22725	53093	A	22853	40	409	WKMRHETRYNVLVSRGVGCH RLFFLNGVFEHLGRAFVQRRKV DTEHQLLRAAVPANQEHRGLT

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ 1D NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22727	53095	A	22858		356	MAGPVKDREAFQRLNFVYQVS LRQGPHGDGARRPRVTAPLPQ AAHCVLAQDPENQALARFYLWH TERTIAKRLVLRRDPSVKEELL CRGCSSLLVPGLTCTPAPRDRC RGQRWTGTDLP
22728	53096	A	22859	2	270	
22729	53097	A	22860	2	364	IFSALPAPRRKVGLNLAPVTEPR DQPWAMIIDVFISRYSGSRGQPR QTLTIKGELKVLMEKELPG\FL QSGKDKDAVDKL\LKDPGRPM GDAQGGTFSEVHPCSVAAIT\SA CHK\YFE\KAGLK
22730	53098	Α	22861	282	469	DCGSELQAPPPPSLPGPLCPSIM PA/PPYASPPMMPMMGPPFRGM MPVGLAPGPMLLQYNTCS
22731	53099	А	22862	7	522	SVWWNSNGLQSMMPKFYCDY COTYULTHDSPSVRKTHCSGRK HKENVKDYYQKWMEEQAQSLI DKTTAAFQQGKIPPTPFSAPPA GAMIPPPPSLPGPPRFGMMPAP HMGGPPMMPMMGPPPPGMMP VGPAFGMRPPMGGHMPMMPG PPMMRPPARPMWPTRPGMTR PDR
22732	53100	Α	22863	3	455	WFLIHDRTHTGERPYKCNECY KAFHRSTCHAHKRTHTGEKP YECNQ*GKAFNSSHSPQIHERT HTGEKPYECKECGKAFKGPSSV RRHERTHSTKKPYECKHCGKA LSYLTSFINHLGMHTGEISHTC KICGKAFYSPSVJQTHE*THT
22733	53101	Α	22864	1	1665	
22734	53102	Α	22865	1	2650	

SEO ID	SEQ ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	eodon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence		09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		<u>. </u>		1.00	laas.	Language outpopping of the
22735	53103	Α	22866	107	2384	MFGVHGTFLGIESRPPLLSLSPS
						LLYTCEMFQDPVACEDVAVNF
	İ					TQEEWALLDISQRKLYREVML
						ETFRNLTSIGKKWSDQNIEYEY
		ŀ				QNPRRSFRSLIEKKVNEIKDDSH
						CGETFTQVPDDRLNFQEKKASP
				ł	İ	EVKSCDSFVCAEVGIGNSSFNM
	l					SIRGDTGHKAYEYQEYGPKPY
						KCQQPKNKKAFRYRPSIRTQER
	i					DHTGEKPYACKECGKTFISHSG
						IRRHMVMHSGDGPYKCKFCGK
		ŀ				AVHCLRLYLIHERTHTGEKPYE
						CKQCGKSFTYSATHRIHERTHT
		1				GEKPYECQQCGKAFHSSSSFQA
						HKRTHTGGKPYECKQCGKSFS
		1				WCHSFQIHERTHTGEKPCECSK
						CNKAFRSYRSYLRHKRSHTGEK
		1		ł		PYQCKECRKAFTYPSSLRRHER
		1				THSAKKPYE/CSFQYHERTHTG
	ļ.					EKPYECKQCGKAFRCAPHLRR
		l				HGRTH/S/AEKPYECKECGKAFR
		ŀ		l		SASHLOIHERTOTHIRIHSGERP
		l			i	YKCKTCGKGFYSPKSFQRHEKT
l		l				HTAEKPYECKOCGKAFSSSSSF
		l		i		PYHERTHTGEKPYECKOCGKA
				ł		FRSASILQMHAGTHPEEKPYEC
		1		l		KQCGKAFRSAPHLRIHGRTHTG
1		l				EKPYECKECGKAFRSAKNLRIH
		1			*	ERTOTHVRMHSVERPYKCKIC
		l				GKGFYSAKSFOIHEKSYTGGKP
ļ		1				YECGLQLPRTAAESVDSEKLQG
1		l		l		PGKQETHVQLLSLYCLRRVLDC
22736	53104	A	22867	1	391	MKEHIERNPVTIKNVVKPSINT
22/30	33104	l^	22007	l'	391	VPLEDMKAHTGEKPYEROKCE
		l		ŀ		LNQRLRDYAFQYHESNSGVVS
ĺ		l				ADSETKSGHSGCLIKEMKNATK
1		l			i	THNGITGGIIFNANHWRETNTIY
1		l				
		l				TRIHI/GEKSCDYKKCGKAFNQ
		l				YSSLRRHES\HTGEKPYERQKC
		l	l	l		ELNQRLRDYAFQYHESNSGVV
		l				SADSETKSGHSGCLIKEMKNAT
		ĺ				KTHNGITGGIIFNANHWRETNTI
		_				YTRIHINKQEKYQATHVSYAAN
22737	53105	С	22868	179	289	
22738	53106	A	22869	3	206	

SEO ID	SEQ ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22739	53107	Α	22870	I	855	MPRAEPRATLGEQEKAGLPLG
						AWRLYLLRHFRKQTELRRSGSF
			i		ì	DVTGALLVAAAVASEAVGSLR
						VAEGGPNTLLLQVLR\SW\PWC
						NKELKTMEERKVKRRSPKSFSA
						HCTQVVNVQKKMSIPVSK\ST\C
						FSNPASQVNFTSRPKVKKEVM
					1	K\EKTKPQGLEEGQRHSSQLPIQ
						HSFLT\DV\$RC\$RKMERGLL\$L\
		1				LNDFHSWKTFKHFGN\ECSI\EQ
						MEHVSGELQEKLARLNLELYG
						ELEELPEDKRKTASDSNLDRLL
						SDLEELNSSIQKLHLA\DAQDVP
22740	53108	Α	22871	67	598	SIRRHCGQRCLVRIRIWPLKLPP
						RLAVREGLRCGTLRDVMPPSM
						SGVVVGPPGP\PPGPCGDTTSGP
						APGTFGQASAPVSPPRRGHGEY
		i			1	LCSSPTPVFFPGLWQPPRAGFTA
						PERSWSCRGPRWQLRAAWAGR
		ı				LSSSPVSALLTFKVIILSTGLLFG
						PLGGWKILSSLPEIFANLAT
22741	53109	Α	22872	1	372	
22742	53110	A	22873	1	3961	FALAPEAGFYSLGQQPSTRISGS
		1			İ	GCVHPVRVWPFSIEPAIGTLNV
						GESMQLEVEFEPQSVGDHSGRL
		i				IVCYDTGEKVFVSLYGAAIDMN
						IRLDKNSLTIEKTYISLANQRTIT
		l			i	IHNRSNIIAHFLWKVFATOOEE
		l				DREKYSKVTAVFSIAASSNSRF
		l	ŀ			ALYESSYNSQRSPSEWACDDLI
		l				KEEKDETDEFFEECITDPLLREH
		1				LSVLSRTFANQRRLVQGDSKLF
		l		1		FNNVFTVEPLGWSERKEMAEIR
			l			DVQKKDTKV
22743	53111	A	22874	1248	1716	WYAEQSAHEATAQLSTPMAIR
122,43	55111	n	22074	1270	17.10	RAVVMPPEORNRTKKEY/SLRI
		l				PFGLPGDRMCFFTHRNDD/PTG
						NEVTPPSLAADLGQQFGGDAT
		[
		l	ŀ			DRQPWMIHQMLIHQGSEQLCF VEHGINMLRLGEILHLAATRHE
				l		
		1		ı		AHFLAFRQFFNHQRSCFALNLA
22744	53112	_	22875	478	663	RQDRITDF
22745	53112	A A	22876	508	884	POPURDOPROPEI PUGUCACO
22/43	22110	<u>ا</u> ^ا	228/0	300	004	RQRHPDQRPGRSLFHGHGMCP
						TTPLQQQVDSQHNGDNNSESA
				I		P*QRRADSIHRSSLHIKKRPPSE
				1		DVSRCFLCIDLAFRQGFTFNKK
						GIRVRIPFGLPGDRMCFFTHRN
		<u> </u>				DDRPAMKLLPFAGNKV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22746	53114	A	22877	li .	506	AFCRFSILVGSAFISETDMCLLC
		1				PFPIGKAISVRLQYFKHETKQSI
l			l			CLGNOKSPLVCPFFGEOGLDSS
		1				SSLDLPSSPNPAPTLCPCVOPPR
						GPVSVSVPVSVMGSRLAPLLSA
						*LFVSVILGRMVILKNPGVLGQ
						ROAGPSPGAPGLPSPSVRAPLG
						HKCSERSPSAT
22747	53115	A	22878	621	877	RRRRRRKQRRKQPWSSPARGA
1						*PGPRRPSLRAAGTRRRRKRRA
		l				SWTRCSNTCPGRCRTCSRSNQA
		ļ				SCPSLLHLLLLLFGSFFSFY
22748	53116	Α	22879	Ī	1339	MPSYKGCGETQRHATNGTAAP
	1	l				PRRSSCRGDAIKLPHAEKKEGR
		1				PPASTSYPLRLGTRSARSSSHC
		1				WCTSWEDPPEATAGEEKRHPK
		1				SPAVCCGEGTVLTASTVTWRA
		ı				EPARGEIWARLCPRDRPGSRAA
		i				AVIWERPCCDRLIADRTGPPNSS
	1	i .			İ	EPHCRAATIPLPTWPLPEGOTLT
	1	l				LKALAKMAAPALLLLALLLPV
		l				GAWPGLPRRPCVHCCRPAWP/L
	ŀ	l				WTLCPGEKGEAGVRGRAGRSG
		l			i	KEGAA\GCPGPAGPQSARRGSV
		l				GPPGAACRRAYAAFSVGRREG
						LHSSDHFOAVPFDTELVNLDGA
		1				FDLAAGRFLCTVPGVYFLSLNV
		1				HTWNYKETYLHIMLNR/QAA/V
	l .	1				VLYAQPSERSVMQAQSLMLLL\
	l	1	1			RRATPVWSVPEELPAVLPGODE
	ŀ		1			LGSLOLHAHHPLADGLVRHEL
	ļ	1				OPGLAGSSRTGHLVFRTDHGLR
	į.	1				OHASHGCPVGPLHVPG
22749	53117	A	22880	532	783	VGSTRGIVEGLIFILGEGGECYY
		1				PRSLFTVANFTVGKRV\KTLNC
		1				PSTDKWINKMCHIHTVEYYSDI
		1			1	KRNEIPMRATCROTLKA
22750	53118	A	22881	275	720	TPIHNCFKENKIPRNPTYKEHEG
-2.00	1	ľ		[1	PLOGELQTTSQGNKRGYKQME
		1				EHSMLMG\GRINIMKMAILPKVI
	1	1				YRFTIPIKLPMTFFTELEKTTSKF
	1	1				IWNOKRARIAKSILSRKNKAG/
	1	1			1	AHHAT*LQTILQGYSNQNSMVL
		1			I	VPKORYRPMEOSPOK
	1			L		VENCKINEVSPOK

SEO ID	SEQ ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence		09/540,217	eodon for peptide sequence	of peptide sequence	deletion, \≔possible nucleotide insertion)
			1	sequence		
22751	53119	Α	22882	1	609	MGRNQSRKAENSKNQSASSTP
		1				KDHSASPATEQSWMGNDFDEL
		l				TELGFRKSVMTNFSQLKEDVRT
	1	1				HHKEAKNLEKRLDK WLTRINSI
		1			l	EETLNDLMELKTMARELRDTC
	1	1			l	TSFSSRFDQVEEGEIQTTIREYY
	İ	l		ŀ		EHLYANKLENLEEMDKFLDTY
	1			l	1	TLPRLNQEEVESLNRPITGSEIE
		1		l		AIINSLPTKKIPGPDRFTA*RRP*
		1	İ	l		MT*WS*KPWHENYVTHAQASV
	Į.	1			İ	ADSIKWKKGKYK\HHQRIL*TP
						LRK*TRKSRRNG*IPGHIHSPKT
	1	1		l		KPGRS*IPE*TNNRF*N*GNN**P
	1	L				TNQKNSRARQIHSQILPEVQRG
22752	53120	A	22883	191	564	PNARKLRTLKKDHSPD**RRKL
	i	ı				RKIK*MQ*KMIKGISQ\RSHRNT
1	1	l		l	l .	NYHQRTL*TPLHK*TGKSRRNG
	i	ı				*IPGHIHPPKSKPGRSQIPEYTNN
		l				KI*N*GSN**PINQKKSRTGRIHS
		L.				QILPEVQRGA
22753	53121	A	22884	337	589	TRRSRRNG*IPGHIHPPKTKPGR
	ľ	l		1		SRIPE*TNNRL*N*GSS**PTNQK
		i		l		KPRTRETDSRILPEVQRGAGTIP
22754	53122	ļ. —	22885	162	539	SETIPNNRKRGTPP TRKSRRNG*IPGHIHPSKTKPGR
22/54	53122	Α	22885	162	339	
1		l				S*IPEETNNKF*N*GSN**PTNQK KPRTROVHSRILPEPISHGMGKS
	İ	l				WKOSL*KPAODKAVPSLIAPIO
1		l		l		HIIGSSGQGNQARERNKGYSNR
		l				KRGSOIVSVCR
22755	53123	A	22886	329	459	NDDTFRKPHRLSPKSP*AADKQ
22/33	33123	r	22000	329	439	LOOSLRIONQCAKLRSIPIHH
22756	53124	A	22887	58	440	GLSROFSPHSVNKAAGEVPTGR
22/30	33124	<u>۱</u> ^	22007	176	1770	GPPQLSKANVARLPLSIPPLWA
1		l			1	GHLWKKGGSPSOGIIDKTLISLG
		i i				QST*GKGQLWVQLQRT*AFLPA
1		l				GSEESSGSPSTA\PNSAKGOTPS
		l			i	SSGSLTRASSSSLYLW
22757	53125	A	22888	219	332	NEGKNLKGSORERNTNYHORT
122.57	155.25	Γ.		[/	J-2	Q*TPLHK*TRKSRRNG
22758	53126	A	22889	1295	1435	ONOROKPHDYLNRCRK/W/HFD
-2,53	[Ι.		1	l	KIOOPSMLKTLNKLGNDGTYL
		1			ŀ	RIIRDI
	1		L			l'ano.

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22759	53127	IA	22890	2	648	YRFSGPTIRSIONHTTHNOGRK
						WHKKGIKKPRAORYESLKGPE
						V*PSGQGVTCLLRADWLSPLLG
1		ĺ				SGTY\VGPKFLRNTRFAKKHNK
				ŀ		KGLKKMQANN\AKAMSARAE
						AIKALVKPKEVKPKIPKGCS/RR
						KLRFDFAYIAHPKLWGRRCFVA
						RFCQGASGCSGPKGKAKVPKA
				i		KVKVS/PKAKAKAKAKDOTKA
				ļ		
1				i	i	QAAAPALV\PAQAPKRTQAPTK ASE
		L.	*****		370	OMYCVFNRNEDACRY/GS/AIG
22760	53128	Α	22891	3	370	
						VLAFLASAFFLVVDAYFPQISN
		l				ATDRKYLVIGDLLFSG\VLASLA
		ł		l		YQRYKAGVDDFIQNYVDPTPD
į.		1				PNTAYASYPGASVDNYQQPPFT
		_				QNAETTEGYQPPPVY
22761	53129	Α	22892	1	297	
22762	53130	Α	22893	3	713	GGGSGGDGDMESGAYGAAKA
		l		İ		GGSFNLRRFLTQPQV\VARAVV
						L\VFALIVFSCIYGEGYSNAHES
1						KQMYCVFNRNEDACRYGSA\IG
		ŀ				VLAFLA\SA\FFLVVDAYFPQISN
		1				ATDRKYL\VIGDLLF\SALW\TFL
1						WFR/VGFCFLTNQWAVTK\PKT
		1				VLVGADSVRAAITFSFFSIFSWG
		1	l			VLASLG\YQRYKAGVDDFIQNY
		1		i		VDPTPDPNTAYASYPGASVDN
				1		YQQPPF\TQNAETTEGYQPPPV
22763	53131	Α	22894	3	219	
22764	53132	A	22895	1445	1777	PVHLGSTCHQCRQKTIDTKTNC
		1		1		RNPRLLGA/CRGQFCGPCLQNR
		ı				YGEEVRDALLDPNWHCP\HCR
		1	1	1	ŀ	GICNCSFCRQRDGRCATGVLVY
		1	1			LAKYHGFGNVHAYLKSLKQEF
		1		1		EMQA
22765	53133	Α	22896	3	337	
22766	53134	Α	22897	425	1209	TMSSTQFNKGPSYQLFAEIKNQ
		1		1		LLFKCDPQKQAALCSWIEGLTR
						LSFSPDFOKSLKEGIVLCTLMN
		ł				KLGSVPKI/NRLMONWHOLENL
		1				SNFIKAMVGYDMNPMDLFEAN
		1			I	DLFESRNMTQNRIL/P/PLDHSTI
					I	SLOMGANKRSSO/VGMTAPGTP
		1			1	OHIYNTKLGAHKCDNSSMSLH
1	1		1		I	MGDTQSANHSVQGFGLGQQIY
1	1	1			I	YPKYCPO/GPAASGAPLDAGNC
1		1		1	1	
1		1				PGPGEAPEDPPCSQEEATTEAPE
		1		1	1	HAGSLHHRPVWVWGIFLYFRL
	L	11		L	L	SFFFL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22767	53135	A	22898	156	559	OHIRETPGAOVHSCGK WNPOA
						LWRTAWQLLT/DTKHTLGKPPS
						/MPHLVIYPLEISC\HTKTCTRML
			l			IASFFIITKTWKRPRCPSVDKWK
		1				NKPWCIHTGLLSSTVPPGEPGC
			İ	i		PIPELSGLLMGLOTTSSRCSVRO
						vw
22768	53136	A	22899	161	359	
22769	53137	A	22900	151	383	
22770	53138	A	22901	238	546	PQEHIMEHLEGVINKPRGRRCR
						PQELQLHYLQKCMDYDGNNLL
						DGFRTLPQPSTHV\HKEEGSEQ
						APLMSEDEL\INIIDGVLRDDDK
						NNDGY\IDYAEFAKSLQ
22771	53139	A	22902	1	819	MAAEDKLLLPQLPELFETSKQL
		l				LDDVEVVTEPAGSRIVQKKVFK
		1				GLDLLEKAAEMLSQL/DLVAEL
1			İ			ELPKTKNSAE\NHTANFSMAYP
						SLIAMASQRQAKRERHKQKEL
						EHRLSAIKSAVETGQADDERVR
			i			EYYLLHLQSLEEIESIDQEIKILR
						ERDSSREASTSNSSRQERPPVKP
		1				FILTRNMAQVKVFGAGYPSLAT
İ						MTVSDWYKQHWKYGALPDQG
	i					IAKATPEEFRKAAQQQEGQEEK
						EEEDDEQTLYRVREWDDWKGI
						HPRDYGSQQSMG
22772	53140	В	22903	238	410	
22773	53141	Α	22904	1	638	VAELELPKTKNSAE\NHTANFS
						MAYPSLIAMASQRQAKRERHK
	į.					QKELEHRLSAIKSAVETGQADD
						ERVREYYLLHLQSLEEIESIDQE
						IKILRERDSSREASTSNSSRQERP
						PVKPFILTRNMAQVKVFGAGYP
						SLATMTVSDWYKQHWKYGAL
			l			PDQGIAKATPEEFRKAAQQQEG
	1		l	l		QEEKEEEDDEQTLYRVREWDD
			l	1	1	WKGIHPRDYGSQQSMG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22774	53142	A	22905	3	1083	RGNSRLRYSHEDELQLPRLPEL
		ľ.		ľ		F\ETGRQLLDEVEVAT\DPAGFR
1						\IVQEK\VFKG\LDLLE\RAAEML
						\SOSTLFGRECRFWKDLSPTHLI
						TFWGP\AFQGA\LPLKQVNPSKR
						LRIILQR\AREHFI\NYLT\QCH\C
1						YHVGQSFEL\PKT\MNNSAENH
						TANSSMAYPMLGAM\ASQRQA
İ						KIQRYKQKKELEHRLSAMKSA
i						VESGQADDERVREYYLLHLQR
						W/VDISLEEIESIDQEIKILRERDS
1		•				SREASTSNSSRQERPPAKPFILT
		l		1		RNMAQAKVFGAGYPSLP/TLW
						TVSDWYEQHRKYGSITGFRGIA
	ŀ					KSAPEGISGKAAQQQEEQEEKE
		l				EEDDEPNNFHRAREW\DDWKD
						TPSLGAYGNRQNMG
22775	53143	Α	22906	30	457	
22776	53144	Α	22907	401	1509	MWKMNCSLLQSKQMCNQLPP
						GSTRDGSWKFRQDPRLGKNA/C
						DCDSCIASEFGVRGYPTIKLLKG
						DLAYNYRGPRTKDDIIEFAHRV
1		l				SGALIRPLPSQQMFEHMQKRHR
1		l				VFFVYVGGESPLKYVTLKEMP
1						AVLVFKDETYFVYDEYEDGDL
						SSWINRERFQNYLAMDGFLLYE
		l				LGDTDLNASSLRERKLILPEDIP
		l				LYVGKLVALAVIDEKNTSVEHT
						SELTVPTVVVLNTSNQQYFLLD
		l				RQIKNVEDMVQFINNILDGTVE
l		l				AQGGDSILQRLKRIVFDAKSTIV
		l				SIFKSSPLMGCFLFGLPLGVISIM
		l				CYGIYTADTDGGYIEERYEVSK
		1				SENENQEQIEESKEQQEPSSGGS VVPTVQEPKDVLEKKKD
22777	53145	A	22908	53	543	LPISSLRTQTLFSSPLFPQHFEHY
22111	33143	^	22900	33	343	LALVSVLPFQLGRGRTAPAKKG
		l				GEK/RKGHSAINEVVIQEYTINI
						HKRIHGVRFKKRAPRALREIPK
		1		1		FAIKEVETLSVRIDSRFNKALCA
1		1		1		KGIRNVPYRIRVOLSRKRNEDE
1		ı		1		DSPNKRYTLVTYVPVSTFKNVQ
		l				TVHVHEN
22778	53 146	A	22909	106	491	IGKGTNELMYGEK/RKGHSAIN
1		ľ		1		EVVIOEYTINIHKRIHGVRFKKR
		l				APRALREIPKFAIKEVETLSVRI
1		l		1		DSRFNKALCAKGIRNVPYRIRV
1		1				OLSRKRNEDEDSPNKRYTLVTY
		1		1		VPVSTFKNVQTVHVHEN
22779	53147	Α	22910	1	321	· · · · · · · · · · · · · · · · · · ·
	F	1			·	

SEO ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
		<u></u>				
22780	53148	Α	22911	313	402	R*GGEKKKGRSAINEVVTREYT
						INIHKRIHGVGFKKRAPRALKEI
		l				RKFAMKEMGTPDVRIDTRLNK
	1	l				AVWAKGIRNVPYRIRVRLSRKR
						NEDEDSPNKLYTLVTYVPVTTF
						KNLQTVNVDEN
22781	53149	Α	22912	1	3801	
22782	53150	A	22913	2	535	GGGMADTVLFEFLHTEMVAEL
	İ					WAHDPDPGPGVSAGLRGEEAG
						ATKGQKMSLSVLEGMGFRVGQ
1	l	l				ALGERLPRETLAFREELDVLKF
l	l	1				LCKDLWVAVFQKQMDSLRTN
		l				HQGTYVL\QDNSFPLLLPMGSG
	ŀ	l				LQV/YLEEAPKFLAFTCGLLRSA
	ĺ					LYTLGIESVVTASVAALPVCKF
						QVVIPKS
22783	53151	Α	22914	1	201	GDLPWEINPLSSCSLLHEKDPPT
ŀ						TSGPQT/DQPKKHHTNFKSGP\T
		l				GNRTVQLTWQPLPEPLELWPK
		\perp				AL
22784	53152	В	22915	226	522	
22785	53153	С	22916	1	705	
22786	53154	С	22917	1	462	
22787	53155	С	22918	I	505	
22788	53156	В	22919	I	1059	
22789	53157	С	22920	I	212	LTSFLHCCLSHHYHGAQIHPRH
22790	53158	A	2292 I	1	212	FLHLICRCHCHQYCHHAYLDPC
		1				PODSHHLHWCWOPLPEPLELRP
			1			RLSD
22791	53159	В	22922	1	864	(KL3D
22791	53160	В	22922	li .	635	
22793	53161	В	22924	1	874	
22794	53162	A	22925	1	566	MPSVRTGDPIILHAHTLNPAVW
22/94	155102	^	22,23	l'	1000	LAWAVEGEGSLTLNIWICPWOI
		l				EPLRHLAYTPGSDATSLTVPTR
	ŀ	l				VISPLCVDSWLFTDPLMKTHAP
		l				ERTFLSFPAAAKROAHLCVLAS
		l				NCDKPMYVKLVKAHCAEHQIN
		1				LIKGDDNKKLGEWEGPCTIDRE
		1				GKPCKVVGCSCAVVKDY/GKE
						SQAKDVIEQYFKCKK
22795	53163	A	22926	59	602	PTAMVEEGIAAGGVMDVNTAV
22193	155105	l^	22,20		I***	QEVLKTALIHDGLARGIREAAK
		1			I	ALDKYVYOSOYCGFLOPEONC
	1	1			1	HPREEGMEFMVLAQKF*MASV
	1	1			1	LPET*EK\RQAHLCVLASNCDEP
1	1				I	MYVKLVEALCAEHQINLIKVD
	1	1			I	DNKKLGEWVGL\CKIDREGKPR
					1	KVVGCSCVVVKDYGKESQAK
					I	DVIEEYFKCKK
					J	PTICETIKOKK

WO 01/075067

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location uf first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon,/=possible nucleotide deletion, \=possible nucleotide insertion)
22796	53164	С	22927	135	293	
22797	53165	В	22928	20	325	
22798	53166	А	22929	677	1031	TPTMCLRSHPLPLALVSVILSFP QPSFSSSLINSSWSSPRSSLSSSSS RSPSSGRVNLHPPQTPSSH*SHL SSRSCCSAPLLTYHSSSLLLVCS VPFPFFPGFPGENPLFVLSGSAP
22799	53167	A	22930	1	1011	MCLRVIFTIGAVSVILSFLSLFFI LDQLQLVIPQSSLSSNLEDPY SSYYAAARRPDSASTEPPPAG KSSAEYRAHFLAVSTEAERIRA DLVLRARPLOPAHPRILUPP AGPFCRMVAHNQVA\ADNAVS TAAEFRRPEP:SPSSSSSSPAAT RAPRICPAVPAPAFGDTHFRTF RSHADVRRITRASALLDALGF YWGPLSVHGAHEPLRAEPV TFVRDSIRQRNCFFALSVKMAS GPTSIRVHFQAGRFHLDOISRES FOLTEELLEHYRGGSRADLLGA TULRQRRVRPLQELCRQRIRRPP VGJRENLLARIPLNPVLRDY\LSS FFFQI
22800	53168	В	22931	180	1466	
22801	53169	Α	22932	I	207	
22802	53170	A	22933	408	3758	CLQOGPGONQHCSHREEAPAV ACQEPFOHESPESVLPLGAPP, SEEDGRQQVGSSRLRHIMAEMI ATEREYIRCLGYVIDNYFPEME RMDLPGGLRGKHHVIFGNLEK LHDFHQQHFLRELERCQHCPLA VGRSFLRHLIGGKQFPESPSGER CTJSSHAPCFWMPHYLPEGPA CPVLRIPCEAFNWKHVECTDPV VSPPKQSPLLFQEEQFGMYVIY SYNKYQSPLLFQEEQFGMYVIY PLGPPTACPALS
22803	53171	A	22934	3	355	EAAELDAMCTATRQWSRAGQ GVTGRPCQGRLPRCGLRSTITP KPIPGGHRKKPLLGAL\GKHTLS YVERWHWEGHSGI*TRDRQNF ERRLTVPPSRQPFAPTDETRCG DDRISFCNYV

SEO ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22804	53172	Α	22935	1928	5312	CLQQGPGGNQHCSHREEAPAV
						AACQEPPGHSEPESVLPLGAPP\
	1	1				SEEDGROOVGSSRLRHIMAEMI
				1		ATEREYIRCLGYVIDNYFPEME
						RMDLPOGLRGKHHVIFGNLEK
		1				LHDFHOOHFLRELERCOHCPLA
				i		VGRSFLRHLIGOKOFPESPSGER
		1				PGLTSSHAPCPWMPHYLPEGPA
		l				CPVLRIPCEAFNWKHVECTDPV
						VSPPKQSPLLFQEEQFGMYVIY
		1				SKNKPOSDALLSSHGNAFFKDK
		1				ORELGDKMDLASY
22805	53173	A	22936	3	306	QREEGERINGERS
22806	53174	A	22937	3	460	LQRQRQHPAAAPAVPVRCFTFC
22000	33174	^	22931	ľ	400	FTDIVIMPKRKSPENTEGKDGS
						KVTKOEPTRRSARLSAKPAPPK
		l		1		PEPKPRKTSAKKEPGAKISRGA
					1	KGKKEEKQEAGKEGTAPSENG
ļ		1				
	ŀ				1	ET\KAEEIHISRSTVNVSTSRGTP
		١.	22222		205	PSTLSVKGQIETVRVKGTEN
22807	53175	A	22938	3	285	SSCPAREQLQRQRQHPAAAPAV
					ŀ	PVRCFTFCFTDIVIMPKRKSPEN
				l		TEGKDGSKVTKQEPTRRSARLS
1		l	l	l		/AVFV*FKLSLQKPAPPKPEPKP
		_				RKTSA
22808	53176	Α	22939	1	413	
22809	53177	Α	22940	2	454	AEVQVEEAGLRRALRVSSCSGR
		l				GSIQRRFYAVPVRCFTFCFTDIV
						IMPKRKSPENTEGIDGS\KVTKQ
						EPTRRSARLSAKPAPPNPEPQPR
						KTSAKKEPG\AKISRGAKGKKE
ŀ	İ	1				EKQEAGK\EGTAPAENDETKAE
						EAQKTESVDNEGRMNCP
22810	53178	В	22941	55	186	
22811	53179	A	22942	2	397	
22812	53180	В	22943	339	470	
22813	53181	Α	22944	1	139	
22814	53182	Α	22945	1	416	MPLLYHLSRCAYVQQCNFANP
				1	l	VVSALCTCLVKPVVEKCTPYA
l	1	1	l		I	VLLEALALRNVRQLEDLVIEAV
	1	1	i		I	YADVLRGSMDQRNQRLEVDYS
	1	1	l	1	I	IGRDIQRQDLSAIARTLQEWG/C
	Ì				1	HQHLLSLRSAASQRQQCGYHK
		1				LHDKQRLLRG
22815	53183	Α	22946	315	570	

SEQ ID NO:	of peptide		SEQ ID NO: in USSN	location of first	codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22816	53184	Α	22947	933	1841	RPAEPDPGSGHPEPKSPTLSAV
						MSAEVKVTGRNQEQFLLLAKS
	1					A\KGAVLATLIHQVLE\APGVY
						VFGELLDMPNVRELAESDFAST
						FRLLTVFAYGTYADYLAEARN
						LPPLTEA\QKNKLSDTFSVV\TL
						AAKVKCIPYAVLLEALAL\R\NV
						RQLEDLVIEAVYADVLRGSLDQ
						RNKRL\EV\DYSIGRDIQRQDLS
	1					AIARTLQEWCVGCEVVLSGIEE
	İ					QVSRANQHKEQQL\GLKHQIES
						EVANLKKTIKVTTAAA\AAATS
			١,			QDPEQHLTELREPAPG\TNQRQ
						P\SKKASKGKGLRGSAKIW\SKS
22817	53185	A.	22948	I	1319	
22818	53186	Α	22949	365	1331	RAGGGAPWRFTGTFATVAPET
	1				1	AADLSLHRHLLSRTTSAWGTA
						CITVPIEPPTWPTWAVPIQWGG
					1	SATCIQEDKLRQGILYGYSQQV
						NQAMDSGLRP\VVDSVNAIRVP
				1		QDYVTQSVPLREMNGSLGVLA
						QQLQNAKLQADAAHSALKQSD
	1					DLKPVFDQAFTKVVTTPADAL
						QPLIPAAQTFTQQLVMVGDYIA
	1				ŀ	QQGTQECGKTFSRASYLVQHS
	İ					RIHIGKKPYECKECGKAFSSGS
	1					YLVQHQRIHTGEKPYECNKCG
			l			KAFTVYGQLIGHQSVHTGEKPF
				i		ECKECGKAFRLNSFLTEHQRVH
						TGEKPFKCKKCGKTFRYSSAP
22819	53187	Α	22950	2	206	GGGQGATSFGRPESGFIRHQPPP
	l					PCRRSSTPTRSKLYTL\SAPE*RS
						VPTSAPGPQDRPPGSVSKKSW
22820	53188	Α	22951	571	1102	MGSVSLGNPQGPFQFTRLPLAN
						VIHPTFFWETGPRGADSLGAKG
		l				RKWQPTFTSRCNLKVYLGRAG
						SQAATGGPSPSLPNLHVKKLRP
						RKAEAWPGRRNTGKVSGR/LA
		l				ALRAPYGESFARAAGPRDAPS/
						QQPRPQPEHAPTLISLGSNFGG
						MVEESWCRMNPDSGRPKEVAP
						WPPPSTK
2282 I	53189	A	22952	1	591	MGPHWKDFKEIDVKKYIDALSI
	1	İ		1		HTYTHTHTHADITDGDHGG
	1		1	I		GALLELEINSHLKVQFRELNITA
		l	1	l		AKETEDGGGGKAIIIFVPFPQLK
				l		SFQKIQVWLRRILPKPTRKSCT\
				1		QNMHWPPRSRTLKAMRDSIPE
						DLVF\PSEIMCMRIHVKLDGSRL
				-		IKAHLDKAQQYHVERNAETFS .
	1	ĺ				GVYMKLTGKDINVEFPEFQL

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hed	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22822	53190	Λ	22953	2	508	APPFELPRAAGIRHEGKFSQEK AMFSSSAKIVKPNGEKPDEFES GISQALLELEMNSDLKAQLREL NITAAKEIEVGGGRKAIIIFVPVP QLKSFQKIQVRLVRELEKKFSG KHVVFIAHRRILPKFTRKSRTK NKQKKPRSRTLTAVHDAIPEDL VFPSEIVGKRNR
22823	53191	Α	22954	145	395	
22824	53192	Α	22955	536	717	
22825	53193	A	22956	1	729	
22826	53194	A	22957	1	1158	
22827	53195	A	22958		2869	MAHLRSCCSSLYQPGHLLPPHN PQDHPLTAPSLTHLTRRAYFTD EKTKGGQSAYAVSLLRECVKL RYSDPTVPLMAAK VCIGSLRW MAAKHLAGVLIHSLSECYWS PLSHPLPEFMGKEESSFATQAL RKPHLYEGDNLYCPKDNIEEAL LLLIESSMLANKLLLLEMEQ QSPGVESGVDRQLSPYSDTNVN AQNAQIVPVAAKSAQNEVPGY PEETEKSRYQSNLITGPQDLRL LLAAPECLPPRMSPPS
22828	53196	В	22959	28	139	
22829	53197	A	22960	60	812	LASPFVPLHRCSAGSOPPCPVP EGLIRIYSMRFCPYSHRTRLVLK AKDIRHEVVNINLRNKPEWYY TKHPFGHIPVLKTSKCQLIVES VIACEYLDDA YPIGRKLFPYDP YERVRQKMILELFGKVPHLPR EFPGSVEDVGENAPNLKAALR\ QEFSNLEEILEYQNTTFFGGTCI SMIDYLETLWFWFERLLECMGY WNCVSHTPALRLWISAMKWD PTVC/ALLMDKEHF\QGFLNPYF QNNPNAFDFGLC
22830	53198	A	22961	2	251	ILLDFKPWDDETDMAQLEACV CSIQLEELV*GASKLVPMGYGI RKLQIQCVVEDDKVGTDLLEEI TKFEEHVQSVDIAAFNKI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		İ		sequence		
	Ļ	<u> </u>				
22831	53199	Α	22962	353	1071	APLWPRAERKFYKQMNGPVAS
				1		ASRQSSAPG/NDHSELVLQIASL
1		l				EVENQSLRGVVQELQQAVSNL
		l		1		EAWLN\IRREKSSPGH/RSTVP
		l				QTQHVSPMRQVERPSPRPPYYD
		l		l		PGQEAEDNEDDDIDLDNKEED
1		1		ľ		KEARRLREERLRLYAKKAKNP
l				l		SLVAKSSILLDIKPWDNKTDMT
	1.0	l		l		QLEACVRSIQLDGLVSGASKLA
		l				SVGYGIRKMQIQCVVEDDKVG
l						TDLLEEEITKFEEHVQSVDIAAL
22832	53200	Α	22963	3	482	PWANEAPAKK/PATPAEG/DAR
		l		l		DDDIDLF\GSDNEE\EDKQAGHS
1				l		CGEER\LR\HYAGEEGPRKPAL/
		1		l		VWPKSLPSLLDVK\PWDDDDGP
		1		l		WPSLEACVR\SIQLDGLVWGGF
1		l	l	l		PSLVPVG\YGIRK\LQ\IQCVV\ED
		l		l		DKVGTDFPWRREIT\KFEEARA
		1				RSVDIASFSTRS
22833	53201	Α	22964	1	2835	
22834	53202	В	22965	571	1337	
22835	53203	Α	22966	Į.	489	
22836	53204	Α	22967	1	618	
22837	53205	Λ	22968	3	5227	SWSRVRPRTRPEGLRQLKMRT
		1				GWATPRRPAGLLMLLFWFFDL
1				l		AEPSGRAVN\DPLTIFHGNTGKC
		1		l		IKPGYGWIVADHCDETEDKLW
		1		l		KWVPQHRLFHLHSQKCLGLDIT
		l				KSVNELRMFSCDSSAMLWWKC
1				ł		EHHSLYGAARYRLALKDGHGT
		ı		l		AISNASDVWKKGGSEESLCDOP
		l		l		YHEIYTRDGNSYGRPCEFPFLID
		1		l		GTWHHDCILDEDHNGPWCATT
		1		1		LNYEYDRKWGICLKPENGCED
						NWEKNEOFGRCYOFNTOT
22838	53206	A	22969	3	453	GOSPILVIYHETKRPSGISERFSG
		1		ľ		STSGNTATLTITGTOAMDEADY
		1		1		YCQAGRDFGGGTKLTVLGQPK
		l	l	[AAPSVTLFPPSSEELQANKATL
	1		l	1		VCLISDFYPGAVTVAWKAHSN
1		l	l	[PVKAGVETTTPSKQSNNKYAA
		ı	1	l		SSYLSLTPEO*KSHRSYSC
22839	53207	A	22970	2	354	30 LEST LEG. VOLVO 19C
22840	53207	IA.	22970	1	315	
22840	22208	IA.	229/1	1 .	313	

SEQ ID NO:	of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X-Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
22841	53209	A	22972	1	712	MAWTLIFITLLTOGTGSWAGA LTOPASVSGSPOGSITISCIGS SDIGAYYFVSWYQQYPGKAPK LLIFDVSGRPSGISSRFSGSKSGN TASLTISGLQAEDEADYYCASY ARSSTVVFGGGTSVTVLGQPKA APSVTLYPPSSEELQANKATLV CLISDFYPGAVVAWKADGSP VNAGVETTTPSKQSNNKYAAS SYYLSLTPEQWKSHRSYSCQV THEGSTVEKTVAPTECS
22842	53210	Α	22973	1	1779	
22843	53211	Α	22975	I	2001	
22844	53212	A	22976	292	1497	TGQLNGFHEAFIEGTTEITSES VGGEHPIDKICDQISDAVLDA HLQQDPDAKVAICETVAKTGMI LLIAGEITSRAAVULPRKVVRE EDIGAGDQGLMFGYATDETEE EDIGAGDQGLMFGYATDETEE EDIGAGDQGLMFGYATDETEE EDIGAGDQGLMFGYATDETEE EDIGAGDQGLMFGYATDETEE EDIGAGDQGLMFGYATDETEE EDIGAGDQGLMFGYATDETEE EDIGAGDQGLMFGYATDETEE EDIGAGDQGLMFGYATDETEE EDIGAGDQGLMFGYATDETEE EDIGAGDQGLMFGYATDETEE EDIGAGDQGLMFGYATDETEE LEDENTROLKEENIKAVVPAKY LDEDITYHLQPSG/RTFVIGGPQ GDAGLTGGRKINDTYGGGQA HGGIGAFSGKDYTKVDRSAAY AARWYAKSLVKGGLCRRVLV OXSYAIGVSHPLSISIFHYGTSQ KSERELL*IVKKNFDLRPGYVR DLDLKKRIYGRAAYGHFGRDS FPWEYPKKLIY

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
W				sequence		
22845	53213	I _A	22977	67	1579	FFVRSLSHTDTNMNGQLNGFH
22843	33213	A	22911	07	1379	EALIEEGTFLFTSKADGEGHPD
	,					
						KICDQISEAVLDAHLHQDPDAK
		1			i	VACETVAKTGMILLAGEITSRA
						AVDYQKVVREAVKHIGYDDSS
						KGFD\YKTCNVLVALEQQSPDI
						AQGVHLDRNEEDIGAGDQGLM
		l			l	FGYATDETEECMPLTIVLAHKL
		l	0			NAKLAELRRNGTLPWLRPDSK
		1			l	TQVTVQYMQDRGAVLPIRVHTI
		ł				VISVQHDEEVCLDEMRDALKE
						KVIKAVVPAKYLDEDTIYHLQP
		1				SG\RFVIGGPQGDAGLTGRIIIVD
						PYDGWGAHEEGAFSGKDYTKV
		1				\SRSAAYGARWVARSLVKGGL
	1	1			ŀ	CRRVLVQVPGPPKKPDEDLYLE
	1				ļ	CEPDPVLALTQTLSFQVLMPSG
					l .	PLPRTSVVPREQQMLPLKDSDL
	1				1	LTQPWYSGNCDRYAVESALLH
						LQKDGAYTVRPSSGPHGSQPFT
						LAVLLRGRVFNIPIRRLDGGRH
		1				YALGREGRNREELFSSVAAMV
						QHFMWHPLPLVDRHSGSRELT
22846	53214	Α	22978	3	325	
22847 22848	53215	A	22979	7	1014 592	LI COODUCE OCTAVOR OF THE
22848	33216	l^	22980	ľ	1392	LLCSGRHSLQGVMGILSFLPVL ATESDWADCKSPQPWGHMLL
					i	WTAVLSLAPVAGTPAAPPKAV
		1				LKLEPQWINVLQEDSVTLTCRG THSPESDSIQWFHNGNLIPTHTQ
	ŀ	1	i			
		1				PSYRFKANNNDSGEYTCQTGQ
		1				TSLSDPVHLTVLSGQWRKAPG
		1				WTWEGQDG*NLLSGRGLQERG
22040	53217	١.	22981	32	1150	VACLLGSIAVSCLSTYQWLFLP
22849	33217	Α	22981	32	1150	EKAVTAVLWAPARLQGVMGIL
		1				SFLPVLATESDWADCKSPQPW GHM\LLW\TAVLFLAPVAGTPA
		l				APPKAVLKL\EPQW\INGGSRED
		1				
						SVTL\TC\RGTHSP\ESDSIQWFH
						NG\NL\IPTHTQPSLQVSRANNN\
		1				DSGE\YTCQTGQTSLSDPVHL\T
		1				VLSEWLVLQTPHLEFQEGET\IV
	1				ĺ	LRCHS\WRDKP\LVKVTFF\HNG
		1				KSKKFSRSDPNFSIPQANHSHSG
		1				DYH\CTG\NIG\YTLLFIPRLLTIT
						VQAPSSSPMGIIVAVVTGTAVA
	1	1				AIVAAVVALIYCRKKRISANPT
				1	Ì	NPDEADKVG\AE\NTITYFTSQW
	1					HPGWLWEEPDDQNRILVSIVLA
1	1	1				LGIWRRNSEREDLVFPGLKFPL
						GEDTGRCCQFPKEKVFLPGVIL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	cudon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
l				sequence		
22850	53218	A	22982	1	471	MKSSFOGHNKEKREKKGPSLFS
22000	55210	ľ	2	ľ		AAYGGGSCLLLSIMAALRPLVK
		l			i	PKIVKKRTKEFIWHQSDQHVKI
l						KCNWRKPRG\QILMPNTGHGSN
į.		l	1			/KKTKHMLPSGFRKFLVPSVKE
Į.		1				LEVLMCNKSPCAEIAHSVSSK\N
			l			KTIVERAAQ/LAVRVTNPNARL
			i			RSEENE
20051	50010	١	22082	102	564	
22851	53219	A	22983	103	364	GIFRGIMAALRPLVKPKIVKKR
						TKKFIRHQSDRYVKIKRNWRKP
			l			RGIDNRVRRRFKGQI\LMPNIGY
1						G\SNKKTK\HMLPKWLSGKFLG
		l			ľ	SHQRSREL/EKLLA*CCNKFFTC
l				1		CPRFVSHCFSFQGTGKGLRGER
		<u>l</u>				SLPNWAFRVTNPNARLRSEENE
22852	53220	Α	22984	3	177	
22853	53221	Α	22985	74	735	RFAGAGAIPEARARPPDVQAAE
		ł				EEKEMDLPDSASRVF/CGDASS
		l l				NMGEHRMNVNAIIL\AQKNML
		1				D\RFEKT\NEMLLNFNNLSSARL
			İ			ASRLSERFLHHTRTLVEMKRDL
		l	l			DSIFRRIRTLKGKLAROHPEAF
						SHIPEASFLEEED\EDPIPTOAPT
		į.	ŀ			TTIATSE\QSTG\SC\DTSPDTV\S
	1	i i	ŀ			PSLEPLASEDLSHVQALASPAI\
ŀ		t		İ		NGPOPDOIDEEMTGRIALPARC
22854	53222	A	22986	3	275	MRTVAEKVDAVRIVHIHSVI\IL
22034	33222	n	22,00	ľ	273	RRSDKRKDRVEISPEOLSAAST
		l	ł			EAERLAELTGRPMRVVGWYHS
l		1				HPHITVWPSHVGKYHGVAMFI
i		1				VFRN
22855	53223	A	22987	1	886	AKMAVOVVOAVOAVHLESDA
22833	33223	^	22901	l [*]	000	FLVCLNHALSTEKEEVMGLCIG
1				l		
1						ELNDDTRSDSKFAYTGTEMRT
						VAEKVDAVRIVHIHSVIILRRSD
1				i		KRKDRVEISPEQLSAASTEAER
l						LAELTGRPMRVVGWYHSHPHI
						TVWPSHVDVRTQAMYQMMDQ
		l		1		GFVGLIFSCFIEDKNTKTGRVLY
		l		1		TCFQSIQAQKSSEYERIEIP\IHIV
1		1	l	1		PHVTIGKVCLESAVELPKILCP/
		l	l	1	l	QEEQDAYRRIHSLTHLDSVTKI
				1		HNGSVFTKNLCSQMSAVSGPLL
				1		QWLEDRLEQNQQHLQEL\HQE
				1		KEELMQELSSLE
						· · · · · · · · · · · · · · · · · · ·

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22856	53224	A	22988	110	988	LKAMSWAKMAVQVVQAVQA
22050	John I	Ι.	22700	1.*		VHLESDAFLVCLNHALSTEKER
				1		VMGLCIGELNDDTSRSDSKFAY
				1		TGTEMRTVAEKVDAVRIVHIHS
l				i		VIIL\RRSDKRKDQVEISPGQLSA
				ı		ASTEAEKLAELTGHPMRVVGW
						YHSHPHITVWPSHVDVRTQAM
				ı		YOMMDOGFVGLIFSCFIEDKNT
				1		RAGRVLYTCFQSIQAQKSSESL
l				1		HGPODFWSSSQHISIEGQKEEER
		1		1		YERIEIPIHIVPHVTIGKACLESA
				1		VELPKILCQEEQDAYWRIHSLT
		1		i		HLDSVTKIHNGSVFTKNLCSQM
1		1				SAVSGPLLQWLEDRLEQNQQH
						LQELQQEKEELMQELSSLE
22857	53225	Α	22989	1	1362	
22858	53226	A	22990	284	542	KLFQKKNKKSSCISDNFFFLLAI
	!	1		I		MVNLLQIVRDHW\VH\VLVPM
		1	i			GFVIGCYLDRKSDERLTAFRNK
		4_				SMLFKSFPHRELQPSEEVTWK
22859	53227	Α	22991	272	1241	FPGGAACDICWVKEENGSLMS
		1				MKMGWKAKSCLPVTVHPRCL
		1				TPYSARLSSTFPL*NSITTGP*QS
				1		PACKRPF*LTTC*GGSRRNSNR
				1		KAA*GPCSPPPPSPPPSPAT\LPR
l		1				GPAGLQPPGVPVLPPLRPRKHY
		1				APGGLP\PPASLLEDDDDTFCTS
		1		1		QAMQPTAPTKLSPPALLPEKDS FSSALDEIEELCPTSTSTEAATA
		1		1		ATDSVKGTSSEAGTOKLDGPOR
		1		1		SRADDSKLMDSLPGNFEITTST
ì		1		1		GFLTDLTLDDILFADIDTSMYDE
				1		DPCTSSSGTASKMAPVSADDLL
				1		KTLAPYSSQPVTPSQPFKMDLT
			ł	1		ELDHIMEVLVGS
22860	53228	A	22992	1	3234	MELKCVCLKNRSPSAEEQPSLL
		1		1		PEPSAQESHGGHPRSALYGRER
i	1	1		1		LTGOPPSSPHPNPAPWASCPAVI
			ł	1		DPVPIYSSVAEAYDSGTEGIPTV
			ł	1		PGLR\HVESGSLRATAQRLPPER
			ł	1		RV/OORTTAPPPROPPRRPLRAS
l		1				LSAAPPEPREPVVAAGPEPGSL
1	1	1	l			QRPARTALPNRPEPQAASPGTL
1	1	1	1	1		VEKYAAPVATQDLLDENPCTS
1		1	1	1		RLGVVDLQAVLTLRKLENLRS
			l	i		GAQTPLEGGQSGRVGPHALRW
		1		1		NRGEEEGERLLC
22861	53229	A	22993	307	1906	
22862	53230	Α	22994	1	1752	
22863	53231	Α	22995	588	704	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nuclcotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22864	53232	A	22996	3219	3407	DQVVWTKRLQDAILVLV*EFRL HSPALRLCVMKQVGMSQGELE
22265	50000	١.	22007		588	WGQFLKLSSRNRKRSGERI
22865	53233	A	22997	1	917	
22867	53235	A	22999	1	1692	
22868	53236	A	23000	1	2266	MKRIRDYYERSHANNLDNLEE
22000	33230	^	23000	l'	2200	MDELLETHNLPGLNHEETENLN
1		1				KPITTSLSPLAGVSSTKKNTKNE
						LSIVLPFTGSILPVGFATRPRGL
i	ŀ			l	i	ASSNPILLHMVAFLFANGTEEL
			i		PVPLMTPTRQKKVIPLIPLMVG	
		1				LGVSASTIALGTGITGISTSVTTF
i						RSLSNDFSASITDISQTLSVLQA
					LVDSSAAVVLQGQLRGLDLLT	
					AEKGGLCIFLNEECCFYLNQSG	
	ĺ	1		i		LVYDNIKKLKDRAQKLANQAR
		1		l		QNLDSMLHGTGMKSDSDQKKS
l		1		i		ENGVTLAPEDTLPFLKCYCSGH
		1				CPDDAINNTCITNGHCFAIIEED
		1				DQGETTLASGCMKYEGSDFQC
l	1					KDSPKAHVRRTIECCRTNLCNQ
		1			l	YLQPTLPPVVIGPFFDGSIRWLV
						LLISMAVCIIA\MIISSSCFCYKH
	l				1	YCKSISSRRRYNRDLEQDEAFIP
						VGESLKDLIDQSQSSGSGSGLPL
		1				LVQRTIAKQIQMVRQVGKGRY
		1				GEVWMGKWRGEKVAVKVFFT
						TEEASWFRETEIYQTVLMRHEN
		1				ILGFIAADIKGTGSWTQLYLITD
ļ						YHENGSLYDFLKCATLDTRALL
	į	ŀ			l	KLAYSAACGLCHLHTEIYGTQG
1	1					KPAIAHRDLKSKNILIKKNGSCC
į.						IADLGLAVKFN\SDTNEVDVP\L
						NTRVGTK\RYMAPEVLDESLNK
1		1		l		NHFQPYIMADIYSFGLIIWEMA
						RRCITGGIVEEYQLPYYNMVPS
						DPSYEDMREVVCVKRLRPIVSN
22869	53237	Α	23001	68	636	GVTSNPKALLGALIAPAFASLPE
						GRGKRIQALNMFRPKNSWRLG
I			1	1	1	LKYPPSSKEKTQVPKTLISGLPE
1	1	1	1	I	I	RKSTTRVGEKLQSAHKMPLSPG
						LLLLLFSGATATA/APLPLEGGP
1	1	1	I	I		TG\RDSEHMQEAAGIRKSSLLTF
I	I	1		1	[LAWWFEWTS\QASAGPLIGKEA
				1		REVARRQEGAPP\QQSARRDRM
						PCRNFFWKTFSSCK
22870	53238	A	23002	1	144	GIRLPSSADWQKTEAQKRREQL
1	1	1		I	1	ILLDELVALVNKRDALVRDLDA
	1	L		I	l	QEKQ

W 0					T	
SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	sequence	noa	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	sequence		02/340,217	sequence	or pepade sequence	l l l l l l l l l l l l l l l l l l l
22871	53239	Α	23003	2	667	RWNSSDPDADRTTLNHADHSS
			1			KIVQHRLLSRQEELKERARVLL
			1			EOARRDAALKAGNKHNTNTAT
						PFCNROLSDOODEERRROLRER
				1		AROLIAEARSGVKMSELPP\YG
						EMAAEKLKERSKASGEONSKL
				ł		VDLKLKKLLEVQPQVANSPSSA
	1					AQKAVTESSEQDMKSGTEDLR
1		ł				
ŀ		ŀ		1		TERLQKTTERFRNPVVFSKDST
		l				VRKTQLQSFSQYIENRPEMKRQ
					<u> </u>	RSIQED
22872	53240	Α	23004	1	1437	
22873	53241	Α	23005	1	2009	MAKPYLYKKYGNQQGVAAIPD
						VPAAWEAEPSPIPSPVLGRKPN
l	1		1			ASQSLLVWCKEVTKNYRGVKI
	i					TNFTTSWRNGLSFCAILHHFRP
	l		1			DLIDYKSLNPQDIKENNKKAYD
	1					GFASIGISRLLEPSDMVLLAIPD
į		1				KLTVMTYLYQIRAHFSGQELN
ŀ	1	1				VVQIEENSSKSTYKVGNYETDT
		1			l	NSSVDOEKFYAELSDLKREPEL
1		l			1	QQPISGAVDFLSQDDSVFVNDS
		1				GVGESESEHOTPDDHLSPSTAS
		1				PYCRRTKSDTEPQKSQQSSGRT
		1				SGSDDPGICSNTDSTQAQVLLG
		ı				KKRLLKAETLELSDLYVSDKK
		ł				KDMSPPFICEETDEOKLOTLDIG
						SNLEKEKLENSRSLECRSDPESP
l			1			IKKTSLSPTSKLGYSYSRDLDLA
						KKKHASLRQTESDPDADRTTL
			1			
						NHADH\SSKIVQQDEERRRQLR
					1	ERARQLIAEARSGVKMSELPSY
	1		1		1	GEMAAEKLKERSKASGDENDN
			1		l .	IEIDTNEEIPEGFVVGGGDELTN
	İ		1			LENDLDTPEQNSKLVDLKLKKL
		I	1		1	LEVQPQVANSPSSAAQKAVTES
	1	1	l		1	SEQDMKSGTEDLRTERLQKTTE
		I	1	l	1	RFRNPVVFSKDSTVRKTQLQSF
		I				SQYIENRPEMKRQRSIQEDTKK
l		I				GNEEKAAITETQRKPSEDEKGF
1	1	I		l	1	KDTSQYVVGELAALENEQKQI
1	1	I	1	i	1	DTRAALVEKRLRYLMDTGQNL
ŀ	1			1		DITUTAL PERIOD OF ALL

SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO: in USSN	Nucleotide location of first		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	sequence	liou	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22874	53242	A	23006	145	3217	LDEYEARLTLANLDDFEEDNEI
						DDENRVNQEEKAAKITELINKI
						NFLDEAEKDLATVNSNPFDDPI
			ŀ			AAELNPFGDPDSEEPITETASPR
			1	1		KTEDSFYNNSYNPFKE\VQTPQ
						YLNPFDEPEAFVTIKDSPPQSTK
				1		RKNIRPVDMSKYLYADSSKTEI
						EELDESNPFYEPKSTPPPNNLV
						PVQELETERRVKRKAPAPPVLS
						PKTGVLNENTVSAGKDLSTSPK
				1		PSPIPSPVLGRKPNASQSLLVW
						KEVTKNYR
22875	53243	Α	23007	1	165	RPAAPRHERRPAADHGHHRPE
]		GRGTGQKACSELPSDEACS/GIA
		_				IG*SLLCSACSVRSRKRQW
22876	53244	В	23008	26	625	
22877	53245	A	23009	325	526	LI CUMPLICITIVINI ICENTA ICENT
22878	53246	A	23010	1946	2224	LLQWRHGLLNPINSPYNSPILPV
						QKPDKPYRLVQHLCLIYQIVLP
						HPMVPNPYTLLSSIPFTTHYSVI
		l				DLKHAFFTIPLYPSSQPLFAFTW
		l				TDPDTHQAQQITLAVLPQCFID
		l				PHYFSQAQISSSSVTYLGIILMK THIGLGAVEOGVVLVGEARAA
			ĺ			QEPMEWVGGSGMVGCRSRAL
		l		[ŀ	RGKAAKARREIERSAVTIVPVL
		l				DENPARHIIPDTTPDHHDCISLIF
		1	l			LTFTPFPHISFFPVPHLEHTWFIE
						GSSTR
22879	53247	A	23011	1	1479	MEENFIRVFFTTTTMLLLIALIK
22017	33247	^	23011	ľ	14/7	OGELCESFDGKSLDGRDERVVI
		1	1			CFDAAAGLHRACSIRGEOONL
		l			1	GTHYFTWQRLFAEVIKLGGLM
	1	1				KPKARNASSLWKEKTKNGFFS
						LETPEERQTSDLHKYQLAFRLF
		l				FIKYKNPAQFMAHLATTLRHF1
		l				ALDPESSHNFONLFSSSHLMHII
						SAPRLLRLYSRFVESPTVTIVPG
						PDFNPASNIIPDTTPDPHDCISLI
			1			HLAFTPFPYISFFPVPHPDHTWF
						IDGSSTRPNCHTPAKAGYAIVS
						TSIIEATALPPSTTSRQAELIALT
						QALTLAKGLRINIYTD/SHILHH
						HAVLWA/E/RGFLTMQGSSIINA
						SLIKTLLQAALLPKEAEVIHCKO
		1				HOKASDPITOGNAYADKVPSL
		1		l		AHAPTLTLAPVLQCPLREVAGT
		1		1		EDDNGKAVLEIPSNIRKSAHVP
		1	1			VOTWKGWDPTNLLGGWFSNL
		l	1			GGFKTLVGTVIFIIGFLLFLPCVI
				1		

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22881	53249	A	23013	27	496	
22882	53250	A	23014	117	210	
22883	53251	A	23015	124	414	
22884	53252	A	23016	3	627	
22885	53253	Α	23017	1	567	
22886	53254	Α	23018	44	551	GGGRSAPLQLGFMSRQAKDD FLRHYTVSDPRTHPKGYTEYKV TAQFISKKDPEDVKEVVVWKR YSDYRKYLHGD/LAYTHRYLFIR RLEEFPAFRAQ\VFGRIVEASV DSRERYRKGAEDLLRFTVHIPAL NNSPQLKEFFRGGEVTRPILEVS RDLHILPPPLIPTPPPG
22887	53255	A	23019	25	488	
22888	53256	c	23020	9	119	
22889	53257	Ā	23021	52	241	
22890	53258	В	23022	4	402	
22891	53259	A	23023	249	984	GSTMYKMNICNKPSNKTAPEK SVWTAPAQPSGPSPELQGQRSR RNGWSWPPPHPLQIVAWLLYLF FAVIGFGILVPLLPHHWVPAQY AVSLGKRRAGPWNPFLSVSAFT ILPQCMGAIFAGHLVVHLTAVS IDPADANVRDKSYAGPLPIFNR SQHAHVIEDILCNLCNVDVSA RSKHCSACNKCVCGFDHHCKW LNNCVGERNYRLFLHSVASALL LNNCVGERNYRLFLHSVASALL CYLLLV/LGGHICLRGVLCOPH ASAHQPTL
22893	53261	A	23025	[11]	1368	OODSGSG ALADGSGI DDDDDGDI
22893	53261	A	23025		1368	QOPSGGALARGSGLPPRFGRL PARSPVOKRKGGWFKGKKKGGWFKGKKKGWFKGKKKGWFKGKKMP PRDLAANRVPTTGYMIELNEQRS QLRATHPDLPFTEIMKILAVQ WAQLSQDKKGRKYAGKALGV TPPPSALTPVALIEEPKQLDIQSL GMHTVTEGLAGRQVILQQVQ GGAWESALTCPWAMAAAAP GPTLTMLQPSISAVGFSQMAMG PHVQGRASKLSIYVYPRRQLST LCMKKEPAAKAAPAAPLSSPGP PALGHAGHGRGLLLERSLTLFH LYELLKYPRASPGLWPPRVWM PHSLTCGTFWVPMIMMRFCAH VASGLLRPPRQVYAHLSPAQG RIPGENGIIVLTSRAAVRIRCSS CSRSLTGNLKKILPRTQSTWSL WRKRDK

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22894	53262	Α	23026	2	405	
22895	53263	Α	23027	1	391	
22896	53264	Α	23028	3	463	
22897	53265	Α	23029	74	466	FAFNMPEPAKSAPAPKKGSKK
		l				AVTKAQKKDGKKRKRSRKESY
		l				SVYVYKVLKQVHPDTGISSKA
		l			0	MGIMNSFVNDIFERIAGEASRL
	İ					AHYNKRSTITSREIQTAVRLLLP
		_				G\EL\AKHAVSEGTK\AVTKYHQ
22898	53266	Α	23030	168	597	KEETDFLFWFCHYCFIMPELAK
			ŀ			SAPAPKKGSKKAVTKAQKKDG
						KKRKRSRKESYSVYVYKVLKQ
1		1				VHVPDTGISSKAMGIMNSFVTTF
		l	1	ļ		FE\RIAGEASRLAHYNKRSTITS
1		l				REIQTAVRLLLPGELAKHA\VSE
		_				GTNA\VTKYTSSK
22899	53267	Α	23031	18	321	FTATWLNQEVGIVGKYRTRYG
			l			VSLQKMVKKIEISQYAKYICSFL
						WQNSKMKRRAVGIWCCG/SCM
				l		KTVKVSSSRNHIVWTYNTTSAV
	*****	Ļ.				TVKSTIRRLKQLKDQ
22900	53268	A	23032	12	445	FTCRSHVSEVEERCVEAPGLLG
						SRSITGFSLHSKQAKRTKKVGIV
	1	1	ļ			\GKYGNPLWGPSLR\KMVKK\IE
ł		1	1			ISQHAQVHFALFCGKTKMKRR
			l			AVGIWHCGSCMKTV\AGGAWT
	ĺ		i			\YNTT\SAVTVKSAIRRLKEFERP
22901	53269	١	23033	3	175	SRTLLYSLRTSLGL
22901	53270	A C	23033	378	434	
22902	53270	A	23034	1	578	MVRCVRLVEAGSVVRYLSTSIC
22903	332/1	A	23033	1	3/8	
					1	RPVVDAGSRALCLQEWADSQQ VKEKQYSSRDVQRAAALNIYRI
						PPSSRKPALCPTPRDRLEYDEDR
						LEHIAYVRARELHTLEVTGLET
					ŀ	VAOSKAHVASLEGLIPEDKVVL
	1					LAGSPLQNEATLGQCGVEALTT
						LEVVGRRLGASL/HTSASKHTM
22904	53272	С	23037	236	379	VRALTYCSSREETFTAI
22904	53272	A	23037	178	534	VVQKTEPRLCFAVSPPFSCSLPL
22905	33213	^	23038	1/0	334	PSLFPHSPLLGRPKRQAKPAAD
İ						EGFWDCSVCTFRNSAEAFKCSI
l						CDVRKGTSTRYLKICVTWVVV
			1			VI*QVFIEHTAICALGVQWYKD
L	L	_				RPGPCSYGA

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
22906	53274	Α	23039	44	1012	VGGGGVQPSMTMGDKKSPTRP
i						KRQAKPAADEGFWDCSVCTFR
	l			ŀ		NSAEAFKCSICD\VRKGTSTRKP
	1					RINSQLVAQQVAQQYATPPPPK
1						KEKKEKVEKAGQKRKPEKRPR
	1					KFSP/SVVTKKNTNKKTKPKSD\
		l				ILKDPPS\EANSIQSANATTKTSE
	1					TNHT\SRPRLEKRGTGGIAQQFG
						QLTVG\NVTVIITDFK\EKTRSSS\
						TSSSTVTS\SAGS\EQQNQRQLR
						GFRGATDKGSLPVPPNPKGRPC
						QAVNDESFLKLAHGNCGKLWN
1						SGVWKFQNLPPAHACLHPWEN
1						LSCGTSTSLVDCCQDNFLLCHG
						HSWPPRDFRHPDDYSGHF
22907	53275	À	23040	1	626	MAGCRNPCKAESHRSKSLLSEA
	l					ESLFREAESQHDRGSKNAGHR
	l					DESVEVWLSLLSSSQTRAEIER
1	l					QSRGKQWLRHLLRDTEGPKET
						VRPELATRAAMDVAEVEFLAE
	1					KELVTIIPNFSLDKSYLIGVDNIP
	İ			i		KADEIRTLVKDIWDTRIAKLRV
	Ì					SADSFVRQQEAHAKPDNLTLM
				Į.		EINPSGTFLTQALNHTYK/LRTN
						LQPSESTQSQDF
22908	53276	Α	23041	1	951	MDAAEVEFLAEKELVTIIPNFSL
						DKIYLIGTWELKYREIRWLSQG
						PHGKPHADPGSSILQNSSGPSG
						VRTNGKSKPWEPFVAEEFAHQ
	l l					FHESVLQSTQKALQKHKGSVA
						VLSAEQNHKVDTSVHYNIPELQ
	1				:	SSSRAPPPQHNGQQEPPTARKG
	1		İ			PPTQELDRDSEQEQEEDDEDGE
	ŀ					DEEEVPKRKWQGIEAVFEAYQ
						EHIEERTLERNRSSTAADKAQL
						LAANRQGHGQPTDPASDNIPKA
	1			Į.		DEIRTLVKDMWDTRIAKLRVS
				l		ADSF\VRQQEAHAKL\DNLT\LM
						EIHTSG\TFLTQALNHMYKLRT
						NLQPLESTQSQDF
22909	53277	Α	23042	1	626	MAGCRNPCKAESHRSKSLLSEA
						ESLFREAESQHDRGSKNAGHR
	1		l			DESVEVWLSLLSSSQTRAEIER
	1					QSRGKQWLRHLLRDTEGPKET
	1		İ			VRPELATRAAMDVAEVEFLAE
	1		1			KELVTIIPNFSLDKSYLIGVDNIP
	1		l			KADEIRTLVKDIWDTRIAKLRV
	1		l			SADSFVRQQEAHAKPDNLTLM
	1		l	l		EINPSGTFLTQALNHTYK/LRTN
1	1		l		1	LQPSESTQSQDF

SEO ID	SEO ID NO:	Ta 1 - 4	SEO ID NO:	N	North did booting of last	Amino acid sequence (X=Unknown.
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1.0.	sequence		09/540.217	codon for pentide	of peptide sequence	deletion, \=possible nucleotide insertion)
	,			sequence		·
		<u> </u>				
22910	53278	Α	23043	2	642	RPRRLLRDAEGPEETVRLWPAA
					İ	RAAMDAAEVEFLAEKELVTIIP
i	1					NFSLDKIYLIGGDLGPFNPGLPV
					1	EVPLWLAINLKQRQKCRLLPPE
						WMDVEKLEKMRDHE\RKEETF
		1				TPMPKPFTTLELT\KLLLNHASD
						NIPKQDEIRTLVKDMWDTRIAK
						LRVSADSF\VRQQEAHAKL\DN
		i				LT\LMEIHTSG\TFLTQALNHMY
						KLRTNLQPLESTQSQDF
22911	53279	Α	23044	3	1038	TSSMAAGCSEAPRPAAG\SDGS
	l	l				LVGQAGVLPCLELPTYAAACA
						LVNSRYSCLVAGPHQRHIALSP
i		1				RYLNRKRTGIREQLDAELLRYS
						ESLLGVPIAYDNIKVVGELGDIY
ł						DDQGHIHLNIEADFVIFCPEPGQ
1	!				İ	KLMGIVNKVSSSHIGCLVHGCF
İ						NASIPKPE\QLSAEQWQTMEIN
						MGDELEFEVFRLDSDAAGVFCI
						RGKLNITSLQFKRSEVSEEVTEN
						GTEEAAKKPKKKKKKKDPETY
	1					EVDSGTTKLADDADDTPMEES
	1					ALQNTNNANGIWEEEPKKKKK
						KKKH\QEVSRTQDPCFSQGQLT
						S\SGYQSDHKKKKKKKKHSEEA
						EFTPPLKCSPKRKGKSNFL
22912	53280	В	23045	82	323	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22913	53281	Α	23046	3	1624	QLEIRAPTA DEIHVTVGEARNLI PMDPNGLSDPYVKLKLIPDPRN LTKQKTRTVKATLNPVWNETF
	l .	l				VFNLKPGDVERRLSVEVWDWD
	ŀ	l				
ļ	1	l		ŀ		RTSRNDFMGAMSFGVSELLKA PVDGWYKLLNQEEGEYYNVPV
		1				ADADNOSLLQKFEACNYPLEL
ŀ		1				YERVRMGPSSSPIPSPSPSPTDPK
į.	1	1				RCFFGASPGRLHISDFSFLMVLG
ŀ		l				KGSFGKVMLAERRGSDELYAI
ŀ		l				
		l l				KILKKDVIVQDDDVDCTLVEKR
•						VLALGGRGPGGRPHFLTQLHST
						FQTPDRLYFVMEYVTGGDLMY
ŀ		l				HIQQLGKFKEPHAAFYAAEIAI
	1					GLFFLHNQGIIYRDLKLDNVML DAEGHIKITDFGMCKENVFPGT
		l				TTRTFCGTPDYIAPEIIAYQPYG KSVDWWSFGVLLYEMLAGQPP
						FDGEDEEELFOAIMEOTVTYPK
		l				SLSREAVAICKGFLTKHPGEAP
	1			İ		GASGP*WGNLTIRAHGFFPLGF
		l				DWERLERLEIPASFSRPRPCGRS
						GENFDKFFTRAAPALTPPDRLV
l	1	l				LASIDQADFQGFTYVNPDFVHP
1		1		į.		DARSPTSPVPVPVM
22914	53282	Λ	23047	h	1522	MLRAPROATROHHFAHIDIVQV
22714	33202	ľ.	23047	1.	1322	TCFRYLRRSALAPAPAMPSPVR
		1				VDRGSFPPVAGGNHTRRRGGTP
		l l		i		OEGVTREALAFFDVVATTGAE
ł	i	1		l		MMWSNFFLQEENRRGAAGRR
İ		1		i		RAHGOGRSGLTPEREGKVKLA
	İ	1		į.		LLLAAWAPRWRCCPWAPSSG
1		1				WTQHLQGQRQRRVRSGPTWG
	i	l				CGRRAPSGCGRRTCPWTGTPA
		l				APRSCPEASERNSEHPALICTLP
		1		İ	İ	QPHMLKRRARRAPGSMAPPLP
		l		l		RPPOAASVPVFPPDWEPLESRA
	į	l				RSKSPLHSQHRPAQETGRHSRP
		l				RTSTRKROOKHSTEKATHTPER
		l				APKPQTRENNPKGRAQGSPAPR
		l				DHNPASRHQKTSNDKRSTHDT
	İ	l				HKTRKDSQTRRKKTRNNEKQQ
1	1	I		1	1	PDR/IGPQIIAYQP/YGKSVDWW
1	1	1			l	SFGVLLYE/MLAGOPP/FDGEDE
I	1	1				EELFQAIMEQ/TVTYPKSLSREA
I	1	1				
I	1	1		1		VAIC/KGCGRSGENFDK/FFTRA
I	1	1				APALTPPDRLV/LASIDQADFQG
1	1	I	1			FTYVNP/DFVHPDARSPTSPVPV
L	L		L	L	L	PASSPTRPQLLENKFGC

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amine acid	
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
22915	53283	A	23048	505	2345	I ISIWGMVMMNPTPSSITLRRMM
22713	33263	<u> </u> ^	23048	1303	2545	SLFPASLTTKYGGFYINSGTLQF
		l		1		RQASESEDDFIKEKKKKSPKKR
		1	1			KLKEGGEKIKKKKKKDDTYDKE
		1		1		KKSKKSKFSKAGFTALNASKEK
		1		1	l	KKKKYSGALSVKEMLKKFQKE
					ŀ	KEAQKKREEEHKPVAVPSAEA
ŀ		1				QGLRELEGASDPLLSLFGSTSD
		1		l	i .	NDLLQAATAMDSLTDLDLEHL
				1	1	LSESPEGSPFRDMDDGSDSLGV
		1		1		GLDOEFROPSSLPEGLPAPLEKR
					l	VKELAQAARAAEGESRQKFFT
						QDINGILLDIEAQTRELSSQVRS
				ı		GVYAYLASFLPCSKDALLKRA
				i		RKLHLYEQGGRLKEPLQKLKE
i		1				AIGRAMPEOMAKYODECOAHT
ŀ		Į.		i		OAKVAKMLEEEKDKEORDRIC
						SDEEEDEEKGGRRIMGPRKKFO
		1				WNDEIRELLCOVVKIKLESODL
				1	İ	ERNNKAQ\AWEDCVKGFLDAE
l	1	1		1		VKPLWPRGWMQARTLFKESRR
l		1		1		GHGHLTSILAKKKVMAPSKIKV
						KESSTKPDKKVSVPSGQIGGPIA
						LPSDHQTGGLSIGASSRELPSQA
	ŀ			1	ł	SGGLANPPPVNLEDSLDEDLIR
				1		NPASSVEAVSKELAALNSRAAG
				1		NSEFTLPAPSKAPAEKVGGVLC
		l		I		TEEKRNFAKPSPSAPPPASSLOS
22916	53284	A	23049	1	379	AYRNFTSASDVWSYGIVMWEV
220		1		ľ		MSYGERPYWDMSNQD\PNTAL
				1		LDPASPEFSAVVSVGDWLOAIK
				1		MDRYKDNFTAAGYTTLEAVVH
				l .		VNQEDLARIGITAITHQNKILSS
		İ				VQAMRTQMQQMHGRMVPV
22917	53285	Α	23050	l	123	
22918	53286	Α	23051	3	221	
22919	53287	С	23052	80	379	
22920	53288	A	23053	71	924	AFSVNSSNSWEKA*SQKCLQV
						NYLSLSSASYLP/CMTCPPSAPL
						NLISNVNETSVNLEWSSPQNTG
						GRQDISYNVVCKKCGAGDPSK
		1				CRPCGSGVHYTPQQNGLKTTK
						VSITDLLAHTNYTFEIWAVNGV
		1		1		SKYNPNPDQSVSVTVTTNQAAF
				1	1	SSIALVQAKEVTRYSVALAWLE
				1	1	PDRPNGVILEYEVKYYEKDQNE
	1			1		RSYRIVRTAARNTDIKGLNPLTS
	l			1	1	YVFHVRARTAAGYGDFSEPLE
				1	1	VTTNTGNKDHPDSVTSEAIFIIQ
1				1	1	LNPITVKSKELSVIFGSADILLQ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22921	53289	Α	23054	1	1102	MRWLLKWERKYTLYSSGVLK
22921	33289	^	23034	[*	1102	HKNLLESLLGQFLGPPAQRFDS
İ						AGGVRPKNDGRFTVIQLVGML
		l				RGIGSGMKYLSDMSYVHRDLA
						ARNILVNSNLVCKVSDFGMSR
	ŀ					VLEDDPEAAYTTRGGKIPIRWT
	ŀ					APEAIAYRKFTSA\SDVWSYGIV
						MWE\VMSYGERPYWDMSNOD
						VIKAIEEGYRLPPPMDCPIALHO
1						LMLDCWOKERSDRPKFGQIVN
					İ	MLDKLIRNPNSLKRTGTESSRP
ļ	l					NTALLDPSSPEFSAVVSVGDWL
l						OAIKMDRYKDNFTAAGYTTLE
						AVVHVNQ/EVSTQRCNTKGDL
	İ				İ	ARIGITAITHQNKILS\SVQAMR
						TQMQQ\MHGRMVPRLSTVLNK
		ļ	20045	140	2220	LKTLEISLPHPCTLIEELHFFNFV
22922	53290	Α	23055	148	3229	NAFRFHLLNPAGESHDRLQDLL
ĺ						SAVRSPTKRQEQRWHRRTMAG
						IFYFALFSCLFGICDAVTGSRVY
						PANEVTLLDSRSVQGELGWIAS
						PLEGGWEEVSIMDEKNTPIRTY
		1				QVCNVMEPSQNNWLRTDWITR
		1				EGAQRVYIEIKFTLRDCNSLPG
l		ł	ŀ		1	VMGTCKETFNLYYYESDNDKE
1		1	Į.			RFIRENQFVKIDTIAADESFTQV
Ì						DIGDRIMKLNTEIRDVGPLSKK
		l	i i			GFYLAFQDVGACIALVSVRVFY
		_				KKCPLTVRNLAQFP
22923	53291	Α	23056	57	450	VGDCLTSRGMSWVQATLLAR/
1		l		1		ALCRAWGGT/CRGALTGTSISQ
1		l		1		VPRLAP\RGLHCSA/ASVSSEQS
1		l				LVPSPPEPRQR/PPGGERDKASF
1	l	l	l	İ	1	LQTVQKFADTSVVDSGHIDFIY
1		l		1		LALRKMREYGVERDLAVYNQL
1		l				LNIFP

SEO ID	SEQ ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	endon for last amino acid	*=Stop eodon, /=:possible nucleotide
	sequence	1	09/540,217	eodon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
22924	53292	-	23057	89	11502	VGDCLTSRGMSWVQATLLARG
22924	33292	A	23037	109	1302	LCRAWGGTCGAAL/SOEPPSLR
						SLASLPRGLHCSAAAHSSEQV\L
					i	
		1				VPSPPEPRQEGPPRLLVPFEGPV LGOGAGLGERGOGRAFL/LDGO
						RNLRTTSVRKRGPH/VDLFYLA
						LRKMRE\YGVERG/DLAVYNQL
ſ						LNIFPKEVFRPRNIIQRIFVHYPR OOECGIAVLEOMENHGVMPNK
1						
						ETEFLLIQIFGRKSYPMLKLVRL
						KLWFPRFMNVNPFPVPRDLPQD
						PVELAMFGLRHMEPDLSARVTI
		l				YQVPLPKDSTGASKIPPKPHIVG
					ľ	HRACMSPHQESKKSRSSKAAL\
						ARHNPS\RPVF\VEGPFSLWLRN
					l	K\CVYY\HILRA\DLLPPEEEGK
					1	WKRTPEEWE\LYYP\RQLDLEY
					i	VRSG\WDNYEFDINEVEEGPVF
				1		AMCMAGAHDQ\ATM\VKWI\Q
į		1		1		GPAGDQPQTLAQNPPWSFRLRR
		1				VQPGELPDILLQGWRSP/HPAR
		_				GPPRKKTTTCKRQQQGPELV
22925	53293	A	23058	1	2517	HITTERIODOL AD ADOCDA LIBORE
22926	53294	Α	23059	1	927	IIVVEFVSRGLARAPSGPAVFSH
						LAMARLPLRGVRWGCLLTAVH
				1		PEPPTACREKQYLINSQCCSLCQ
1					1	PGQKLVSDCTEFTETECLPCGE
I						SEFLDTWNRETHCHQHKYCDP
1						NLGLRVQQKGTSETDTICTCEE
						GWHCTSEACESCVLHRSCSPGF
					i	GVKQIATGVSDTICEPCPVGFFS
						NVSSAFEKCHPWTSCETKDL\V
ĺ						VQQAGTNKTDVVCGPQDRLRA
						L\VVIPIIFGILFAILL\VLVFIKKV
						AKKPTNKAPHPK\Q\EPQEINFP
İ				1		DDLPG/SSNTAAPVAGRIFTWG
		_		L	100	GQPV\TQEDGQRESHFSCRERQ
22927	53295	В	23060	24	439	
22928	53296	A	23061	1	582	
22929	53297	С	23062	1 0 40	2607	+ TOUTTO FOR AUGUST CONTE
22930	53298	Α	23063	248	430	ATNYTREQDSMWNCVR*GRTT
1		1		1	I	ATTHSVTQRTDERQRRADTAL
00001	50000	 _	22064	200		TLADYRNTSVTGVFAGLF
2293 I	53299	A	23064	776 284	1426	VEDI I DOOVEEDOVEDOV 1 431
22932	53300	Α	23065	284	598	VFRLLRGQKSEPGKSRPKAL*N
1	1	1		1	I	KPANTPVQSHISLYAAGIINSDS
1		1		1		KVR*SCRQSSARQSGSYL*TSPR
	1	1		I	i	SPQNRQQTGEITATVIAFGRTRR
22224		L.	02055		2500	TAPDITPHPASAR
22933	53301	A	23066	3139	3792	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22934	53302	Ā	23067	2	830	GVPNMA APODVHVRICNOEIV
1		l				KFDLEVKALIODIRDCSGPLSAL
		l				TELNTKVKEKFOOLRHRIOPVL
		l				YORAFIWTASTFFFKLTYSLTDF
		1	ŀ			SSTOHDFNSPTTPVTFSDL\EQL
		l	ŀ			AKEODKESENOLLLOEVENHK
		1				KOMLSNOASWRKANLTCKIAI
						DNLEKAELLOGGDLLRORKTT
						KESLAQTSSTITESLMGISRMM
1		ı				AQQVQQSEEAMQSLVTSSRTIL
		1		ľ		DANEEFKSMSGTIQLGRKLITK
		1				YNR\RELTDKLLIFLALRLFLAT
						VLYIVKKRLFPFL
22935	53303	A	23068	1231	1458	
22936	53304	В	23069	1	2133	
22937	53305	Α	23070	722	957	VFRLLRGQKSEPGQKQAKSTLE
		i	1			QTSEYPSNRGIPIIRQSECSIRTPL
		l				PFVRALRD*VRGCGCAPLSSVD
		_				SFFCCTEAL
22938	53306	Α	23071	2	1137	VPNMAAPQDVHVRICNQEIVKF
						DL\EVKAPIQDIRD\CSGPLS\AL
						TELNPKVKEKFQQLRHRPTGP/
			l			VEQLAKEQDKESEKQLLLQ\EV
						ENHKK\QMLQQFRPSWEGKLN
						SSPGKICNSDNLEKAELLQGGD
						LLRQRKTTKE\SLAQTSSTIT\ES
						LMG/ISGRMMAQQVQQSEEAM
						QSLVTSSRTILDANEEFKSMSGT
			İ			\IQLGRKLITKYNRRELTDKLLIF
		1				LALALFLATV\LYIVKKRLFPIF
	l					VRFPKVPVLAPFNSLFSGICPWF
						PEAPRLPKAEAHTPLRFAPQLP
		l				CRLGWEHQCPPLFLQSSWLVK
	1					GEQSPWEEGPGIGMPPLGTYEP
						PPSHHCALWEAAAAWSTLKSE
						PPRTRMPLLSPAPIVPLNARSGG
		ļ.—			***	GKREER
22939	53307	Α	23072	64	665	RVRGGEAELAMAAAEEE\SGAP
						NVPNRERGGAGRPKTPFECNIC
	1					LETAREAVVSVCGHLYCWPCL
						HQWLETR\PERQECP\VCKAGIS
	İ					REKVVP\LYGRGSQEAPRIPELK
						TPPPPPRGQR\PAPESRGGF\QPF
		1	1			GDTGGFHF\SFGVGAFPF\GFFT
		1				TVF\NAHEPFRRGTGVDLG\QG
		L		L	L	HPSLQLGKEFPLPGFSAIFFLFW

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nuclcotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22940	53308	A	23073	1	416	MQRRRRRRLLLLPPRRQTDRAL PPPPSPGSMPPPLPPPP/PPARGG GLRPGDGLYPK/PSPARRGSGA
		1		l		RTSWCTMCAPPSTSAPRASRRP
		1		l		RPSSCATRPPTSKPPPA\GHAPH
		l				PPPRDLGGGLDSGVQSDGVLQ
				ŀ	1	HLORPGH
22941	53309	Α	23074	1	1713	
22942	53310	Α	23075	1305	1538	LKVKRNCTSS*SPTIGHD*QSNL
						SLPPSDFAKPPFPFSLLRCSKFR
						KLKAAYRKRKKGHKSSLSLSV
		1				KINKTAAANK
22943	53311	Α	23076	49	355	
22944	53312	В	23077	253	471	
22945	53313	A	23078	1267	1497	LKVKRNCTSS*SPTIGHD*QSNL
						SLPPSDFAKPPFPFSLLRCSKFR
İ						KLKAAYRKRKKGHKSSLSLSV
		l				KINKTAAANK
22946	53314	Α	23079	3	1237	HEVAGAVPGAIMDEDYYGSAA
		1			1	EWGDEADGGQQEDDSGEGED
1					ŀ	DAEVQQECLHKFSTRDYIMEPS
		l				IFNTLKRYFQAGGSPENVIQLLS
		i			ł	ENYTAVAQ\AVNLLAE\WLIQT
1		1				GVEPVQVQDTVEYHLKSLLIKH
						FDPRIADSMFTEEGETPAWLE\Q
				i		MIAHTTWRDLFYKLAEAHPDC
						LMLNFTVKLISDAGYQGEITSV
l		1				STACQQLEVFSRVLRTSLATILD
ŀ						GGEENLEKNLPEFAKMVCHGE
1						HTYLFAQAMMSVLAQEEQGGS
}						AVRRIDQEVQRFAQEKTVLTQE
		1				AIITVKGVSLSSYLEGLMASTIS
						SNASKTIVAAMSNLVPPVELAN
						PENQFRVDYILSVMNVPDFDFP
						PEFYEHAKALWEDEGVRACYE
						RSNEYQLIDCAQYIFVLEMQVP
			L			CCEEAQSHPGQHGGVQVGRN
22947	53315	A	23080	1358	1589	LKVKRNCTSS*SPTIGHD*QSNL
1		1		1		SLPPSDFAKPPFPFSLLRCSKFR
		l	İ			KLKAAYRKRKKGHKSSLSLSV
	L					KINKTAAANK
22948	53316	A	23081	33	458	GITGSTDASGVTMKFNPFVFSD
						RRKNRKRHVNAPSHVRRKIMS
			ĺ	l		SPLSKELRQKYNVRSMPIRKDD
	l			1		EVQVVRGHYKGQQIGKVVHV
1	1	1				YRKKYVIYIERVQREKANG/DK
			ĺ			DRKKILERKAKSRQVGKEKGK
L	L	<u>L</u>				YKEELIGKMQEYIEP
22949	53317	Α	23082	1	576	
22950	53318	В	23083	37	276	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22951	53319	Α	23084	124	549	NNSLSSPVQADVYFPRLIVPFCG
						HIKGGMRPGKKVGVMGIVDLN
	1	l				PKSFAINLTCGDSEDPPADGAIE
	l	l				LKAVFTDRQLLRNSCISGERGE
		l				EQVSNPLLSIHSTTMPFRVEILC
		1				EH\HCFR\VFVDGHQLFDFYHPH
		1				SNRYLAI
22952	53320	Α	23085	1	215	QGFIKPPLKEENFTLPPPGKFGP
						PKGSLNSPPS/PSSP*SPGGGILLL
		1				VYYLYRALFQKGCKEAAWLEY
						PLT
22953	53321	Α	23086	533	983	PWGPGCNKRWPCLAKDPPPTP
		1				QSSGVGQPFGGAEPAGAYIRTG
		1	İ			PHLGPPPPPPPDSSTNLHSPNPS
	İ	ı				DDGADTPLAQSDEEEERGDGG
	1					AEP\G\PAASSGPLAYRPDPAGY
		l				SPTSRPPGPSPEPLG/AEDPDSLL
	1	1				ASRAPRGGNGWWDCGTLS
22954	53322	Α	23087	1422	1670	NRYIDKTGWAWWLTPVIRHFR
	ŀ	1				RPRWVDCLRLGVREQPGQHSKI
						KSLQKNAK\ISQAWWHAPVDP
		l				ATWKADAGGSLEPGSLRLQ
22955	53323	Α	23088	2	663	VLLDERSAALDGAKRDGTLAL
			i			AAGALCREARAAQVFFLKGGY
						EAFSASCPELCSKQ/INVSANCP
		1				NHFEGHYQYKSILCGMTTHKA
		1				DISSWFNEAIDFIDSIKNAGGRV
						FVHCQAGISRSATICLAYLMRT
						NRVKLDEAFEFVKQRRSIISPNF
		1		i		SFMGQLLQLESQVLAPHCSAEA
						GSPAMAVLDRGTSTTTVFNFPV
				1		SIPDHSTNSALSYLQSLITTSSHC
22956	53324	Α	23089	3	206	

3019

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22957	53325	Α	23090	982	2252	LFLEKRRPALGPENRQRAPQGR
						STLGALRPAMVMEVGTL\DAG
		İ				GLRAL\LGERAAQLPACLDCRS
						FF\AFNAGHIAGSVNVRFSTIVR\
	ł					RRAKG\AIGALE\HIVPNAELR\G
						RLLAGAYHAVVLLDERSAG\LD
	İ	i				GAKRDGTLALAAGRA/LCREA
		i				RAAQVFF/LSKGGYEA\FSASCP\
						ELCSKQSTPMGLS\LSLSTSVPD
		l			1	SAESGC\SSCSTP\LYD\QGGPVE
	ŀ	ı				ILPFLYLGSAYHASRKD\MLDA
		l				LGHRLPWINVLSQIVPNHF\EGH
						\YQYKSIPVEDNHKADISFWFN\
	İ					EAIDFIDSIKNAGGRVFVHCQA
		l				GISRSAT\ICLAYLMRTNRVKLD
	i i					EA\FEFVKQRRSIISP\NFQLHGA
		ŀ				SLLQFEFPRMLGPALFRAEAGK
						PRQMAVL\DRGT\STTTVFNFPV
		l				SIPVHSTNSALSVPFRAPLTDLS
	1	1				QLVERPTGR
22958	53326	Α	23091	13	287	
22959	53327	Α	23092	1	492	
22960	53328	Α	23093	1	636	
22961	53329	A	23094	1	506	PTRTWTRGRIPRLSAQPSRGAR
		l				GTMADPRVRQIKIKTGV\VKRL
		ŀ		1		V\KEKVLYEKEPKQ\QEEKIEKM
		l				RAEDGENYDIKKQAEIL\QES\R
		1	1			MMIP\D\CQRR\LEAAYLDLQRI
i						LENEKDL/EKEAEEYKEAR\LVL
	Į.	l				DSVKLESLKLFSYGVVFALKPR
		 -				GPILQSIIFDHVLCVQV
22962	53330	A	23095	1	786	ULC A FUNCCI PROPET A FEBUR
22963	53331	Α	23096	183	1059	VLGAFIWQGLDPSPETAFTDHR
ļ.		1		1		ATEWGTNDQSLCLNSP*\DAPS
		l				AKHEYQASVSGSVSSKPEIDLQ
		l				LYDEVRDQDVPVKRNFARIVV
		l		1		NVSDTNDHAPWFTASSYKGRV
	ŀ	l				YESAAVGSVVLQVTALDKDKG
	İ	l				KNAEVLYSIESGSHPPYKTAAD
		l				VLLKAPPPGWRPTNTKPVHSTE
		l		1		AQPTQNQRTQQKHNQGASRSP
	1	l	1	1		RHSFATSTRAGAGICGCKTRRW
	1	l		j		MTRWMEESRGSLQTLPSTRLEP
		1				GSSAAWLDPEEQAQSLQFSSQE
				1		APFLGEGGENHIKGVPHGTKES OOOPLNHRSSL
22964	53332	Α-	23097	1	1224	QQQFLININGSL
22964	53333	_	23097	1	660	
22963	22333	Α	23098	Li.	1000	l

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				Sequence		
22966	53334	Α	23099	129	595	LFPRLLSCLTTPPHCSFSICFVIC
1						SRTLILKGSSLMYVFCLPNTAIV
						MALSPRGWRSKFGMPWHSKPT
1	İ	ł				WIMTL*FPLEKTGALSLTSDTS
		l				MFTVVVLTLGGLPLSEA*T/GG
						NYDSHFDVDKGTGTIIVAKPLD
						AEQKSNYNLTVEATDGTTTILT
22967	53335	Α	23100	1	1782	MAARIEVIHELSIRLLLMKTDL
						DAATAALLNVQPAKSGDNTSG
						APFPGKISQTLLKDPHNGYHVV
		1				FALAGSPKDADDTSIYMFYQK
	İ					VGDNSIDSWKNAGRVFKDSDK
		l				FDANDPILKDQTQEWSGSATFT
						SDGKIRLFYTDYSGKHYGKQSL
		l				TTAQVNVSKSDDTLKINGVED
	İ					HKTIFDGDGKTYQNVQQFIDEG
						NYTSGDNHTLRDPHYVEDKGH
						KYLVFEANTGTENGYQGEESLI
						NKAYYGGGTNFFRKESQKLQQ
						SAKKRDAELANGALGIIELNND
						YTFEKSNEAADHFKHGGGATIK
					İ	GRVADNGKPQLSSLTYIDIRVIE
						ESIYPPAILPLEIFITASGEEYSGO
						VIGKIHATDQDVYDTLTYSLDP
						QMDNLFSVSSTGGKLIAHKKLE
						IGQYLLNVSVTDGKFTTVADIT
						VHIRQVTQEMLNHTIAIRFANL
			1			TPEEFVGDYWRNFQRALR\NIL
						GVRRNDIQIVSLQSSEPHPHLDV
1						LLFVEKPGSAQISTKQLLHKINS
1			1			SVTDIEEIIGVRILNVFQKLCAG
1						LDCPWKFCDEKVSVDESVMST
1		1				HSTARLSFVTPRHHRAARQQ\R
						GPPLHNPADT*TTARKYNCDKR
22968	53336	Α	23101	184	13958	AMGRHLALLLLLLLLFQHFGDS
		1				DGSQRLEQTPLQFTHLEYNVTV
j		1				QENSAAKTYVGHPVKMGVYIT
j		l				HPA WEVRYKIVSGDSENLFKAE
		1				EYILGDFCFLRIRTKGGNTAILN
		1				REVKDHYTLIVKALEKNTNVE
1	1	1				ARTKVRVQVLDTNDLRPLFSPT
1		1				SYSVSLPENTAIRTSIARVSATD
1		ı	ĺ	1	I	ADIGTNGEFYYSFKDRTDMFAI
	l	1	1			HPTSGVIVLTGRLDYLETKLYE
	l	1		1		MEILAADRGMKLYGSSGISSMA
	l	1	1			KLTVHIEQANECA
22969	53337	С	23102	56	331	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
22970	53338	A	23103	2249	2680	GEFLKTREFSIYFLEKYTTFPGE NLEHGFWDSSTSPPVLLHPGIPA SRI/PVSSIFTXVPPKKSATKP/PV SIWGAPPLGPPGFWGCCPTGEQ RGVCHPSEPHPPVVDFRKKPPF LCVSLSLINIFKSNKFLLTTCNL VIFYTCSL
22971	53339	A	23104	739	1672	COEAASEFGCPLHTPAMFLIRIL GGWLPRPWGRRKPMRPDPPYP EPRRVDSSSENSGSDWDSAPET MEDVGHPKTKDSGALRVSRAA SEPSKEPEPQVEQLGSKRMDSLK WDQPISSTQESGRLEAGGASPK LRWDHVDSGGTRRPGVSPEGG LKGVPGFGAPLEKPGRBEKLLG WLRGEFGAPSRYLGGFEECLG WLRGEFGAPSRYLGGFEECLGS STINLTHLELELLASALLAGLELASAL LSFLAALIGHAVLSLTTAHL LFSLAALIGHAVLSLTTAHL HFACLFGLLQALVLAVSLREPN GDEAATDWESEGLEREGEEQR GDPGKGL

3022

SEQ ID	SEQ ID NO:					Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	sequence	or pepude sequence	detection, v-possible nucleotide insertion)
	<u> </u>	<u> </u>				
22972	53340	Α	23105	122	2437	RLELKLPLWTMRFTEMELLGH
				ĺ		LSIFKITGKQKQLTRTAPGPSGQ
	ł			l		LRSNQTTGNSLEPSGQPRSSQTT
			1			GNSLEPSGQPRSSQTTGNSLEPS
			i			GQPRSSQTTGNSLEPSGQPRSSQ
		1				TTGNSLEPSGQPRSSQTTGNSLE
	Ì	1				PSGQPRSSQTTGDSLEPSGQLRS
		1				NQTTGDSVQPNNREQPGTVRS
		1				APLQPNNREQPGTVRSAPLQPN
l		1				NRGQRPTKQQGTTWNRQVSSA
		1		1		PTKQQGTAR\PSGQPRSNQTTG
1	1					NSLEPSSQLRSNQTTGDSVQPN
İ						NREQPGTVRSAPLQPNNRGQRP
				1		TKQQGTAR\PSGQPCSNQTTGN
						SLEPSSQLRSNQTTGDSVQPNN
						REQPGTIRSAPLQPNNRGQRWD
						RQVSSAPTKQGTALGPSG\SSAP
		1				TKQQGTALEPSSQLRSNQTTGN
l	1			1		SLEPSGQPRSNQTTGNSREPSSQ
			i .			LCSNQTIGNSQDRQVSSAP/IQT
1	1	1				TGNSQDRQVSPAPTKQQGTAG
		1		ļ		NRQVSST/PIQTGNSVQPNNREQ
}						PGTVSGAM/VWVATPVKGPAR
				1		VETSAHKVSAAGLPKKKT/QPG
	1		1	1		VTGGAPRLTLNTVLPKHSRTPS
				i		RHEAGSTQALYHNGQAQYALS
		1				FPHEAAPMAAFYVVTGIEVTTA
		1				LTLEEAPCTVEITTALTLEEPPC
						TVFWNKTERNTRTLVQDKGSN
			ļ			TKGFVQTATCCFRADCKASRPS
		1				PHPPSPHREWTSDLRHKQVSKF
		_				PGAHRLPGHPELPIQEAWDRGE
22973	53341	Α	23106	605	813	LFVFINFRENPALLRWAYAR\TI
						NVYP\NFRPTPKNSLMGALCGF
						GPLIFIYYIIKTERVSIQTRCLVF
22974	53342	Α	23108	273	936	AYSNKVPSWNQKA\PSPTKPAG
1						TLIFDFPAVKTVRNKFVFFINYP
					1	VSGYTEDLKKFLKPYTLEEQKN
						LTVCPDGALFEQKGPVYVACQ
						SPISLLQACRGMNDLDFGYSQR
1		1				NPCILVKMNRIIGLKSEGVPRID
		1	l			CVSKNEDIPNVAVYPHNGIIDL
						KYFPYYGEKLHVGYLQPLVAV
1		1	1	1		QVSFAPNDTGKEVTVECKIDGS
		_				ANLKSQDDRDKFLGRVMFKIT
22975	53343	Α	23109	104	478	ETEVGGLVEASMSHNCTTALQ
		1	1	1		PGQHSQKENLQNKPIPGREKY*
		1		1	1	VSIEFELIYFP/YYCQVGYLQPL
1		1	1	1	1	VAVQVSFAPNNTGKEVTVECKI
		1		1		\DGSANLKSQ\DDRDKFLG\RV
L			L		L	MFQNHSTVHSMSRVSSTE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
22976	53344	Α	23110	3	903	LLSPVRAHTMTKNEKKSLNQSL
						AE\WKLFIYNPTTGEFLGRTAAS
						WGLILLFYLVFYGFL\AALFSFT
						MWVMLQTLNDEVPKYRDQIPS
						PGLMVFPKP\VTALEYTFSRSDP
						TSYAGYIEDLKKFLKPYT\LEEQ
				ļ		KNLTV\CPDG\ALFEQKGPVYV
			İ			ACQFP\ISLLQACSGMNDPDFG
			i		l	YSQGNPCILVKLNRIIG\LKPEG
					l	VPRIDLCFKRMKDIPNVAVYS\
						HNGMIDLKYFPYYGKKTAMF/
						VGIYQPIGLLVQVSFAP*QPLGK
						EVTVE/CAKIDGSSQPKKVQDD\
						RDKVFGDEFMFQIHSTVH
22977	53345	Α	23111	467	931	SRDSRRGTPLAGGRGRDFAGQ
						MGHRVQGRHGSRGRLPSGDAR
				ŀ		VNAERAEPVGAGAGGPGGSS
		ł				VRGFCGRSWSEA/RGRRRAGG
						GVH\IEP\RYRQFP\QLTQIPRCSQ
					1	SEF\FSGLMWFWIF\WRFWHDP
						K\KVLGHFPV/YPDPSQWTD\EE
						L\GIPPDD
22978	53346	Α	23112	1	390	VTIPTSLAELQGLPHIRVFLDNN
					l	PWVCDCHMADMVTWLKETEV
						VQGKDRLTCAYPEKMRNRVLL
		1			1	ELNSADLDCDPILPPSLQTSYVF
		1			1	LGIG*ALIGAIFLHYRNNGPLTIS
		1			1	SEGGLASLLCKDSRGVPLL

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
22979	53347	Α	23113	3	1653	PQRIQRGVPTRGEEVAPGRRQE
		1				EEREQERGAERPDPPCVLSGGK
l		l				GRGNRPLREAPGGEGVSQVPA
						AAPLPRFHERKVFFFQTLPPAR
						ALRAQPPEPSERAPSQPSSGETR
l				1		AAMPGGCSRGPAAGDGRLRLA
						RLALVLLGWVSSSSPTSSASSFS
l						SSAPFLASAVSAQPPLPDQCPAL
		l				/CACRSEAARTVKCV\NRNLTE
l						VPTDLPAYVRNLFLTGNQLAVL
						PAGAFARRPPLAELAALNLSGS
						RLDEVRAGAFEHLPSLRQLDLS
			ŀ			HNPLADLSPFAFSGSNASVSAPS
						PLVELILNHIVPPEDERQNRSFE
						GMVVAALLAGRALQGLRRLEL
			Ì		1	ASNHFLYLPRDVLAQLPSLRHL
		ļ				DLSNNSLVSLTYVSFRNLTHLE
ł						SLHLEDNALKVLHNGTLAELQ
						GLPHIRVFLDNNPWVCDCHMA
						DMVTWLKETEVVQGKDRLTC
				1	1	AYPEKMRNRVLLELNSADLDC
		l				DPILPPSLQTSYVFLGIVLALIGA
						IFLLVLYLNRKGIKKWMHNIRD
					1	ACRDHMEGYHYRYEINADPRL
						TNLSSNSGCLEEILEDRPRTTLH
22980	53348	A	23114	288	877	ASRLSLPTRTSMRSSSSTSPSSSS
		l				SSSSLCRFLLNSRVTDAAFNFLL
						VWYYCTLTIRESILINNGSRWG
						RAGPEGGEHGMGCQGPSGKGR
				l		EQGAGG*WWVFCSRV\KGWW
						VFHHYVSTFLSGVMLTWPDGL
						MYQKFRNQFLSFSMYQSFVQF
						LQYYYQSGCLYRLRALGERHT
			ĺ			MDLTVGESGVALASGHSPFLPS

3025

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*-Stop codon, /-possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22981	53349	Α	23115	1	1273	MAPLHSILGNRVGLCQASVRA
						CPGATVSSQCRASLRSRLAWRG
	ł				1	SELPANRGRTWPGPGGLRLOA
	1					APRLAYLGRATLLVLTPYYDLP
					ĺ	PNVRFTSMPASPHTYFLVLGAPI
					i	
	1			i		FLLLRSTWSYYHACRFEYATPN
						DRNTTYLAISGPTRRTYDVATF
						AYKDEYEKFKLYLTIILILISFTC
		1				RFLLNSRVTDAAFNFLLVWYY
						CTLTIRESILINNGSRPDGLMYQ
	1		i			KFRNQFLSFSMYQSFVQFL\QY
	1					YYQSGCL\YRLRALG\ERHTM\D
						LTVEGFOSW/MWRGLTFLLPFL
						FLW/YNSGQLFNALTLFKPGPR
						DPQLQGVGRCLCCGFPFLL\LFL
	l					
						GNFFTTLRVVHHKFHSQRHGS
	ļ					KKDLRLGLSPCRPRRGFLVPVC
						FWEGDWEGAPSSVRVSGGLFY
						SPLGFMGRCGPGKEDPGPSCPQ
22982	53350	Α	23116	238	398	
22983	53351	A	23117	1	386	
22984	53352	Α	23118	13	580	
22985	53353	В	23119	39	1171	
22986	53354	A	23120	381	795	PPSSKAPSSKGPAKIVCIYWALI
						LOSNEVTITEDKFNNLIKAAAV
		l				TVEPFWPSFFAKALASVNIGSLL
						CNAGVG/SMAPAACTAQAGGP
ı						ATSTTAASASFRNHNPRIAPKFR
						GVLAHEQYLSERATWVWAPNP
		_				NPKSLF
22987	53355	Α	23121	1	960	
22988	53356	A	23122	2	3952	
22989	53357	Α	23123	1058	1494	KSVRLVVN/YLRTQKAVVRVSP
	1	ı	1			EVPLQNILPVICAKCEVSPEHVV
1		l	l			LLRDNIAGEELELSKSLNELGIK
		1	l			ELYAWDNRRVPSAVPMOGPWP
		1				AEHFLOVNEOPPAGSPSPCPSSS
1		ı				SSSSGLRGGMOTANGPACALTP
1		l	l			
22222	50050	ļ. —	00101	105	200	TCAQTDCPCPTSH
22990	53358	Α	23124	195	398	TDSSSALQPQHRGRPALRQRTP
						PRRSP/SPGLSPPRGASPAS\SGA
1		1	l			PARGVAFGGDEPPIGAEAMYSK
			l			P
22991	53359	Α	23125	2	155	
22992	53360	Α	23126	1	339	
						

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22993	53361	A	23127	150	1017	LGRWRODOVPHGLECRSOSEW
22993	33301	l^	23127	150	1017	SCTDSROHSSS\YSDSRROEIGK
1					1	MRYVSVRDFKGKVLIDIREYW
						MDLKRLPRKPRTGEAGSGSLSS
				ŀ		PGADTPVEAAVTWRLGLGSVD
						VGSFRAWQRGKSAAVGHTVSP
				ŀ		RWPRWQGGDSSALSMASLSTL
					ŀ	TCSWGLRYSALSPTTSAQREEG
						SALYRHLSPPSLRFYPTVKWAT
					ľ	LAITRPRPPAPRSHSHHPQPRAL
						CPVWVLFQHPKTHGENLGDRW
						KEGSALVSSKGTPAKLSSGRIR
1				ŀ		NALQNTLGEVSVARIQTHRGIF
						VETTEARL
22994	53362	A	23128	I	584	APIECGGIPSLPVLCQSRANDQE
ł		l				GVRLLEPSEAMPKSKELVSSSSS
						GSDSDSEVDKKLKRKKASLL/P
		1				ENPVKK\QKTGETS\RALSSF*TE
1						PASSRDDNMFQIGKM\RYVSV\
		1				RDFKGKSA*FDIREYWMDPEGE
						MKPERKGISLNP\QQWEPSLKE
		l				QISGPLIDASKKTCKISEPILIKPC
		ı				TVPVVLICLVTVAVCF
22995	53363	Α	23129	186	687	LLWARGLGRAKSAVPTVSTML
į		1				GLPWKGGLSWALLLLLLGSQIL
ļ		l				LIYAWHFHEQRDCDEHNVMAR
İ						YLPATVEFAVHTFNQQSKDYY
ĺ						AYRLGHIFNSWKEQVESKTV\F
İ						SMELLLGRTRVWEYLKTTIDNC
						HFOESTELNNTFTCFFTISTRPW
ĺ		ı				MTOFSLLNKTCLEGFH
22996	53364	A	23130	232	402	LFKAWECFFLIFLOVVITRLKLD
		l				KDRKKILERKAKSRQVGKEKG
		ı				KYKEETI\EKMOE
22997	53365	A	23131	637	758	NVEEKNRKKILERKAKSCOI/K
		1				KEKGEYKEETIGTMRECGL
L	1			L	l	TELECOL TELECTION TO THE COLD

SEO ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	coden for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
						L
22998	53366	Α	23132	1	1095	IVFLLYLETCLEVMDDKPNPEA
		l		l	1	LSDSSERLFSFGVIADVQFADLE
			i	l		DGFNFQGTRRRYYRHSLLHLO
		1	1	i		GAIEDWNNESSMPCCVLOLGDI
		1		1		IDGYNAQYNASKKSLELVLW\Q
		l		į.		FKRRKGPVHHTWGNHEFYNFS
		l				REYLTHSKLNTKFLEDQIVHHP
				į .		ETMPSEDYYAYHFVPFPKFWFI
		l				LLDAYDLS/ALGAWDQSSPKYG
		1		İ		GSVWKILEGSHNP\NTGNWNSS
1		l				SRDFSEA/RSLFQFNGRVSAQEQ
1		l				
		l				LNWLNEVLTFSDTNQEKVVIVS
		l		ĺ		HLP\LYPGPPFDNVCLAWNYRD
		l	ļ.			ALAVIWSHECV\GVSLLGHTHD
						GGYSEDPFGVYHG\NLEGVIET
ŀ		ł				APDSQRFGTVHVYPDKMMLKG
						RGRVPDRIYELQGKEEPSHC
22999	53367	С	23133	235	382	
23000	53368	В	23134	63	293	
23001	53369	Α	23135	160	330	
23002	53370	В	23136	1	870	
23003	53371	Α	23137	2	1013	
23004	53372	Α	23138	I	1552	
23005	53373	В	23139	120	1034	
23006	53374	Α	23140	1	584	RTRGRRSLAFLLLAERGAAAA
		l				VLWSGCRRFRARLGCLPGGLR
İ		l				VLVQTGHRSLTSCIDPSMGLNE
ľ		ı	1			EQKEFQNAAFDFAAREMAPNM
1		l				AEWDOKELFPVDVMRKAAOL
ŧ		l				GFGGVYIQTDVGGSGHVRLDT
1		l				SVIFEALA\TGCTSTTAY\IRHPO
1		ı				CPOSOHSFVOEGGTHYKAGHL
1		l				CCROWGPRACLPPGRPCHRGA
23007	53375	A	23141	2	412	CONQ II GITATODITI GITA GITACOT
23008	53376	Ā	23142	47	487	
23009	53377	A	23143	77	459	GWKGIEHTLGRCLEAOGVINY
2300)	33377	ľ.	23143	l''	1737	RMLLSWRRWLVAHPVF*SFSCE
	i					VGPVGQS*RMTMLHCRHSDPV
i						AKRLSCRRGDFASVVVLCPPPO
1						RLRTLKHFLFFGGLPLSIDFVMT
	44470	١		ļ.———	***	VDFCCHLHLTDFSSNFGF
23010	53378	A	23144	1	304	KVAGSPSPTPAGQQVLDVNSCL
			1			DSSVLDSSFLTFSGLHAE/AEGS
1		l	l			TGGPAALAYLGLKTLCTSQQAF
1		l	l	1		AGQMKSDLFLDDSKSDLEEPW
						ELRKEGAPRSRDRE
23011	53379	Α	23145	1	207	
23012	53380	Α	23146	1	3009	
23013	53381	A	23147	3	298	
23014	53382	Α	23148	1	636	
23015	53383	Α	23149	1	795	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23016	53384	Α	23150	125	1527	SSPROOTOLARVHP/DGRAO/LA
						PSCLPGLGLCPLPVCPPVCNCPA
						CPCHEHCGEAP*GVA\AKRGAL
						RVHPHPSCGGPRAMTVTLHGQ
		ŀ			•	DVLMLPGDLVGLQHDAGPGAL
						LHCSPAPGHPGPOAPYLSANAS
		ŀ				SWLPHLPAQLKGTWACPACAL
						RLLAATEQLTVLLGLRPNLGLR
						LPGRYEVRAEVANGVSRHNLS
İ		l				CSFDVVSPVAGLRVIYPAPRDG
						RLYVPTNGSALVLOVDSGANA
		ĺ				TATARWPGGSVSARFENVCPA
						LVATEVPGCPWETNDTLFSVVA
						LPWLSEGEHVVDVVVENSASR
1						ANLSLRVTAEEPICGLRATPSPE
					İ	ARVLQGVLVVLLAGSSGYLVG
						FKFLESHGSDSGSANSFHCLISQ
ŀ						NEFKTLLPDLTRAPRYSPVVEA
İ						GSDMVFRWTINDKQSLTFQNV
					l	VFNVIYQSAAVFKLSLSADSAP
						RPRWRLQSPRVGRLSPDPGFFG
						SPGPCQPRLQAFLI
23017	53385	A	23151	648	1387	ASTOSLLAWCPAWLSIPSTWPS
23017	33363	^	23131	048	1307	SFSSGCPG/CKAOPVYEVSRSSY
						VYLEGRCLNCSSGSKRGGYTFT
						LT\VLGRSGEEGCASIPLSPNRP
				l		PLGGSCRLFPLGAVHALTTKVH
1				l		FECMGWHDAEDAGAPLVYAL
1						LLORCROGHCEEFCVYKGSLSG
						YGAVLPPGFRPQFEVGLAVVV
						QDQLGAAVVALNRSLAITLPEP
						NGSAMGLTVWLHGLTASVLPG
				l		LLRQADPQLVIEYSLALVTVLN
						ETYTESL
23018	53386	A	23152	2	654	DKHIWLSIWDRPPRSCFTRIQRA
23018	33300	^	23132	 	054	TCCVLLICLFLGANAVWYGAV
			1	I		GDSAYSTGHVSRLSPLSVDTV\
1				į.	l	AVGLVSSVVVYPVYLAILFLFW
				I		MSRSKV\AGSPSPTPAGOOVLDI
				l		DSCLDSSVLDSSFLTFSGLHAEV
1					i	INTLADHQHRGTDFGGSPSVLII
	-			1		TVSLRSYKFAISLCTTYLWVINT
				1		LADHRHRGTDFGGSPWLLIITV
	L		L		L	FLRSYKFAISLCTTYLC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
		1		sequence		
23019	53387	A	23153	2	4072	SRDPAGSYHLNLSSHFRWSALQ
23019	33361	l^	23133	ľ	4072	VSVGLYTSLCOYFSEEDMVWR
		l			\	TEGLLPLEETSPRQAVCLTRHL
		l			ì	TAFGASLFVPPSHVRFVFPEPTA
		1				DVNYIVMLTCAVCLVTYMVM
		l				
						AAILHKLDQLDASRGRAIPFCG
		1				QRGRFKYEILVKTGWGRGSGT
						TAHVGIMLYGVDSRSGHRHLD
		1				GDRAFHRNSLDIFRIATPH\SLG
						SVWKIRV\WHDNKGLSLALGFL
						Q\HVIVRDLQTA\QSTFFLVNDW
		<u> </u>				LSVETEANGGLVEKE
23020	53388	Α	23154	1	10922	MPPAAPARLALALGLGLWLGA
		1				LAGGPGRGCGPCEPPCLCGPAP
					Ì	GAACRVNCSGRGLRTLGPALRI
						PADATALTAMAKADGDVKKY
I						GSKRGALALTHGHSLLRDVSH
ł			1			NLLRALDVGLLANLSALAELDI
1						SNNKISTLEEGIFANLFNLSEINL
			1			SGNPFECDCGLAWLPRWAEEQ
			1			QVRVVQPEAATCAGPGSLAGQ
						PLLGIPLLDSGCGATLVGPHGPL
						ASGQLAAFHIAAPLPVTATRWD
						FGDGSAEVDAAGPAAS
23021	53389	A	23155	3887	14184	HVLCAWVTYMVMAAILHKLD
					ŀ	QLDASRGCAIPFCGQRGRFKYE
						ILVKTGWGRGSGTTAHVGIHA
				1		VWGGQPERPPAPGTATEPSTAT
						VWDIFQIRHPAQPGVACWKIRV
						FGTTTKGSALPRFPCSTSSSGTC
			- 4			RRHAAPSSWSMTGFRWRRRPT
						GAWWRRRCWPRVRPRSVFPLR
						HAALLRFRRLLGAELQRGFFDK
						HIWLSIWDRPPRSYFTHIQRATC
					Į.	CVLLICLFLGANAVWYGAVGD
					1	SAYSTGRVSRLNPLSVY
23022	53390	Α	23156	1	442	GGASGMGGIT\ASTVNQ\SLLSP
	1	1			1	LVLEVDP\NIQAVRTQEKEQIK\
1	1		1			TLNNKFASFIDKVRFLEQQNKM
1	1		1	l	l	LETKWSLLQQQKTARSNMDN
1	1		1			MFESYINNLRRQLETLGQEKLK
						LEAEFTTWGPPLPMPPATKQFN
1						CFFFLVQNKTSASSANVK
23023	5339I	A	23157	I	497	,
23024	53392	C	23158	208	422	
		_			·	L

NO:	SEO ID	SEO ID NO.	Met	SEO ID NO-	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
23025 53393 A 23159 81 1339 ELSSPRSSFSAPSRISAWFGPF TPASTMSIRGTQKSYKGSTSG RAFKQAAS*QRSGPGFRISSS SRVGSNFRGGLGGGGGGG A MGGITAVTVNQ)SRALLSPL EVDPNIQAVROPECEQIKTJ KFASIFIDEVGFLEQONKML KWSLLDQQKTAKRSIMDING ESYINNLRRQLETLAQEKLK AELGINMQGLVEDFKNNYED NKSTETENEFYLIKKDVDEA NEXTETENEFYLIKKDVDA NEXTET							
23025 53393 A 23159 B 1339 ELSSPRSSFAPSRISAWFGFF TPASTMSIRGTQKSYKGSTX RAFKQAAS*QRSGPGFRISSS SRVGSNNRGGLGGYGYGGA MGGITAVTVNQISRALLSPL EVDPNIQAVRTQEKEQIKTLI KFASIFIDEVGFLEQQNKML. KWSLLDQQKTAVRSNMDIN ESYINNLRRQLETAQEKKLI AELGNMQGLVEDFKNNYED NKSTETENEFVLIKKDVEA NELYEEIRELGGISDTSVV DNSRSLDTDSIJAEVKTEISEN RNISWLQAETEGLAGGRISLI GIADAEQRRELVIKDANAKL LEAALQQTKKLLEMENMSII KTTSGVAGGVSLAVGGLTG LSYGLSSSSRSGAGSSSFSRT RAMVVKKIKTROCKLLTLS LSYGLSSSSRSGAGSSSFSRT RAMVVKKIKTROCKLTLS LSYGLSSSSRSGAGSSSFSRT RAMVVKKIKTROCKLTLS LSYGLSSSSRSGAGSSSFSRT RAMVVKKIKTROCKLTLS LSYGLSSSSRSGAGSSFSRT RAMVVKKIKTROCKLTLS LSYGLSSSSRSGAGSSFSRT RAMVVKKIKTROCKLTLS LSYGLSSSSRSGAGSSFSRT RAMVVKKIKTROCKLTLS LSYGLSSSSRSGAGSSFSRT RAMVVKKIKTROCKLTLS LSYGLSSSSRSGAGSSFSRT RAMVVKKIKTROCKLTLS LSYGLSSSSRSGAGSSFSRT RAMVVKKIKTROCKLTLS LSYGLSSSSRSGAGSSFSRT RAMVVKKIKTROCKLTLS LSYGLSSSSRSGAGSSFSRT RAMVVKKIKTROCKLTLS LSYGLSSSSRSGAGSSFSRT RAMVVKKIKTROCKLTLS LSYGLSSSSRSGAGSSFSRT RAMVVKKIKTROCKLTLS LSYGLSSSSRSGAGSSFSRT RAMVVKKIKTROCKLTLS LSYGLSSSSRSGAGSSFSRT RAMVVKKIKTROCKLTLS LSYGLSSSSRSGAGSSFSRT RAMVVKKIKTROCKLTLS LSYGLSSSSRSGAGSSFSRT RAMVVKKIKTROCKLTLS LSYGLSSSSRSGAGSSFSRT RAMVVKKIKTROCKLT LSYGLSSSSRSGAGSSFSRT RAMVVKKIKTROCKLT LSYGLSSSSRSGAGSSFSRT RAMVVKKIKTROCKLT LSYGLSSSSRSGAGSSFSRT RAMVVKKIKTROCKLT LSYGLSSSSRSGAGSSFSRT RAMVVKKIKTROCKLT LSYGLSSSSRSGAGSSFSRT RAMVVKKITROCKT LSYGLSSSSRSGAGSSFSRT RAMVVKKIKTROCKT LSYGLSSSSRSGAGSSFSRT LSYGLSSSSRSGAGSSFSRT RAMVVKKIKTROCKT LSYGLSSSSRSGAGSSFSRT LSYGLSSSSRSGAGSSFSRT LSYGLSSSSRSGAGSSFSRT RAMVVKKIKTROCKT LSYGLSSSSRSGAGSSSFSRT RAMVVKKIKTROCKT LSYGLSSSSRSGAGSSSFSRT LSYGLSSSSRSGAGSSSFSRT LSYGLSSSSRSGAGSSSFSRT LSYGLSSSSRSGAGSSSSRST LSYGLSSSRSGAGSSSRST LSYGLSSSRSGAGSSSRST LSYGLSSSRSGAGSSSRST LSYGLSSSRSGAGSSSRST LSYGLSSSRSGAGSSSRST LSYGLSSSRSGAGSSSRST LSYGLSSSRSGAGSSSRST LSYGLSSSRSGAGSSSRST LSYGLSSSRSGAGSSSRST LSYGLSSSRSGAGSSSRST LSYGLSSSRSGAGSSS	t		1		codon for peptide		deletion, \-possible nucleotide insertion)
### PASTMSIRGTQKSYKQSTSK RAFKQAAS*QRSGPGFISSS SRVGSSNFRGGLGGGGGGGG MGGITAYTVNQ)SRALLSPL EVDPNIQAYROPEKEQIKTJ KFASFIDEVGFLEQONIML KWSLLDQKTAYRSNMIDN ESYINNLRRQLETLAQELKI, AELGNMQGLVEDFKNNYED NKSTETENEFYLIKKDVDEA NEXTETENEFYLIKKDVDAA NEXTETENEFYLIKKDVAA NEXTETENEFYLIKKDVAA NEXTETENEFYLIKKOM NEXTETENEFYLIKKOM NEXTETENEFYLIKKO					sequence		
### PASTMSIRGTQKSYKQSTSK RAFKQAAS*QRSGPGFISSS SRVGSSNFRGGLGGGGGGGG MGGITAYTVNQ)SRALLSPL EVDPNIQAYROPEKEQIKTJ KFASFIDEVGFLEQONIML KWSLLDQKTAYRSNMIDN ESYINNLRRQLETLAQELKI, AELGNMQGLVEDFKNNYED NKSTETENEFYLIKKDVDEA NEXTETENEFYLIKKDVDAA NEXTETENEFYLIKKDVAA NEXTETENEFYLIKKDVAA NEXTETENEFYLIKKOM NEXTETENEFYLIKKOM NEXTETENEFYLIKKO	23025	53393	IΔ	23159	81	1339	EL SSPRSSESAPSRISAWEGPPAS
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FNQPWRKAGGSICKGGSQKC ISAPSAECPPCFVMGCEPMKI DKQAFKNISNASCTNQLL+H AKG*FHDNFGNSLEGLHDHK MAITANPRKTVD/GPSRGKL\							VKAENGKLVINGNPITI\FQE\RD
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ISAPSACPPCFVMGCEPMK# DKQAFKNISNASCTNQLL*H AKG*FHDNFGNSLEGLHDH# MAITANPRKTVDIOPSRGKL LMGRGALKOJIPBASTGVAK, GKVIPELNGKLTGMAFRVPT VSVVDLTCRLERPAKYDDIK VVKQASEGPLKGLIGVTTERIJE VSDFNSDTHSSTFDAGAIG, NDHFVKLISWYDMEFGYSNI VJLMVAHMGFQGS 23029 33397 A 23163 160 236 23030 53398 A 23164 931 1124	1		1				FNOPWRKAGGSICKGGSQKGSS
AKĞPHDNEGNSLEGLİIDIK MAITANPRKTVDIĞPSRĞKLÜ LMĞRĞALÜNIPASTĞYAK, GKVIPELNÖKLTĞMAFREVPT VSVVDLTCKLEKPAKYDDIK VVKQASEĞPLKĞILĞYTEHQ VSSDFNSDTHSSTFDAĞAĞI, NDHFVKLISWYDNEFĞYSNI VDLM'AHMĞFQĞS	1		1				ISAPSAECPPCFVMGCEPMKKY
AKĞPHDNEGNSLEGLİIDIK MAITANPRKTVDIĞPSRĞKLÜ LMĞRĞALÜNIPASTĞYAK, GKVIPELNÖKLTĞMAFREVPT VSVVDLTCKLEKPAKYDDIK VVKQASEĞPLKĞILĞYTEHQ VSSDFNSDTHSSTFDAĞAĞI, NDHFVKLISWYDNEFĞYSNI VDLM'AHMĞFQĞS	1		1				DKQAFKI\ISN\ASCTNQLL*HPL
MAJTANPRKTVD/GPSRGKL LMGGAL/QNIIPASTG/AAK, GKVIPELNGKLTGMAFRVPT VSVVDLTCRLEKPAKYDDIK VVKQASEGPLKGILGYTEHQ VSSDFNSDTHSSTFDAGAIG, NDHFVKLISWYUDNEFGYSNI VIDLM/AHMGFQGS 23029 33397 A 23163 160 236 2360 33398 A 23164 931 1124			1				
LMGRGAL\QNIIPASTG\AAK. GKVIPELNGKLTGMAFRVPT VSVVDLTCRLERPAKYDDIK VVKQASEGPLKGILGYTEHQ VSSDTSDTHSSTFDAGA\GI. NDHFVKLISWYDNEFGYSNI VDLMAHMGFQGS 23029 53397 A 23163 160 236 236 2303 33398 A 23164 931 1124			1				MAITANPRKTVD\GPSRGKLWP
GKVIPELNGKLTGMARRVPT VSVVDLTCRLEKPAKYDDIK VVKQASEGPLKGILGYTEHQ VSSDPNSDTHSSTFDAGAGI, NDHFVKLISWYDDMEGYSNI VDLM\AHMGPQGS 23029 33397 A 23163 160 236 236 23030 33398 A 23164 931 1124			1				LMGRGAL\QNIIPASTG\AAKAV
VSVVDLTCRLEKPAKYDDIK VVKQASEGPLKGILGYTEHQ VSSDFNSDTHASTFDAGA/GI, NDHFVKLISWY\DNEFGYSNI V\DLM\AHMGFQGS 23029 53397 A 23163 160 236 23030 53398 A 23164 931 1124	1	1	1		l		GKVIPELNGKLTGMAFRVPTAN
VSSDFNSDTHSSTFDAGA\GI, NDHFVKLISWYIDNEFGYSNI V\DLM\AHMGFQGS 23029 53397 A 23163 160 236 23030 53398 A 23164 931 1124					ŀ		VSVVDLTCRLEKPAKYDDIKK
VSSDFNSDTHSSTFDAGA\GI, NDHFVKLISWYIDNEFGYSNI V\DLM\AHMGFQGS 23029 53397 A 23163 160 236 23030 53398 A 23164 931 1124	1		l				
NDHFVKLISWY\DNEFGYSNI V\DLM\AHMGFQGS 23029 53397 A 23163 160 236 23030 33398 A 23164 931 1124		1	1		1		VSSDFNSDTHSSTFDAGA\GIAL\
VIDLM\AHMGFQGS 23029 53397 A 23163 160 236 23030 53398 A 23164 931 1124	1		1		1		NDHFVKLISWY\DNEFGYSNRV
23029 53397 A 23163 160 236 23030 53398 A 23164 931 1124							
23030 53398 A 23164 931 1124	23029	53397	A	23163	160	236	
בטכבון בטכבון בטכבון בטכבון בטכבון בטכבון בטכבון	23031	53399	Α	23165	132	303	

	ptide ho		location of first	codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
		09/540,217	codon for peptide	of peptide sequence	deletion, \"possible nucleotide insertion)
			sequence		
23032 534	00 4	23166	120	1171	
		23167	1008	1262	HELOCSLLPVRHATAGHTNGQ
23033 534	01 A	23107	1008	1202	NRLHWGHVAHSNYGAALKTN
			1		GVEVR*FTAAIARCDGNPYESF
		20160	700	11/0	RKDVRADFQTRIDAIVRCLPK
23034 534	02 A	23168	732	1162	RCHEKGRKGHQAPTGSTRPDG
					TLSACRSCQQGQGMPASAGVP
- 1			1		ARQMGSTGNLPHLV*PYFLGLS
				<u> </u>	SKAQHTG*LHKKR
23035 534		23169	2	1093	
23036 534		23170	236	3084	
23037 534					LRDGRGSMSSSSPNSASGSAICP
23038 534	06 A	23172	305	522	
	i				ASSVMPETVSVSRVMPNVTKP
					A/MTAKQTRIRNQMVAITYIES
				***	DAWNRSYQ
23039 534		23173	177	386	C + 1/4 DE GUDY A UDATOC A A 4 CODO
23040 534	08 A	23174	3	200	GAVAPMHPLAHPVPGAAAQPP PA*QPATPCHQPA*GQPAPPDV
		20105	 	101	PGEPSAGRSLHHAHICLAPNEQ
23041 534		23175	171	535	LIPTORIOTEL TOLICUPE OFFICE
23042 534	10 A	23176	190	505	VPTDRIRTRLTSVCKRRCEVHQ
	- 1				LEDRPFEVYQVA/PRRD*SFETS
	- 1		1		VWNHMRKQ/PPLDVRQLAS*N
		1			VLHRQLDIMDSAVASMYSIHLS
			0.10	1070	IKMGIGFSGVTTVVKGTPK
23043 534		23177	349 518	1272	VSSSIIMFGPLSQFPRQQVGSSV
23044 534	12 A	23178	218	1451	KRPSNVVCPGLMFSFFSSAA*IP
					**PLI*OPGV
					TSPDVLRAGPRVPSVOGTSPSO
23045 534	13 A	23179	2436	2894	
	- 1				SCPGQRPHGPNSKAYGR/LSVM
	- 1			ŀ	AQYYCNVIPVLE/VPPSAFTPPP
					KVDSAVV/RLVPHATMPHPVK
				i .	DVRVLSRITTEAFNQRRKTIRNS
				į .	LGNLFSVEVLTGMGIDPAMRA
					ENISVAQYCQMANYLAENAPL
23046 534	14 A	23180	1846	2557	IPVSERQGSDWRFFSVYVVNKA
					GGLIYQLDSYAPTGLRLEKTF\S
		1			YPLDLLLKLHDERVLVAFGQR
				1	GRHPEWGHAVL\AINGMDVNG
					RYTAD\GKKVLEYLG\NP\ANYP
			ı		V\SYLFAQPRL\TSNDKLMLAL
			1		MFHSLFAIGSQACLP\EQGKPQG
	1		ı		IENAWKTDTFKLH\CYQTLT\GI
		1			KFVVLADPRQAGIDSLLRKIYEI
			1	1	YSDFALKNPFYSLEMPIRCELFD
					QNLKLALEVAEKAGTFGPGS
23047 534		23181	34	225	
	16 B	23182	105	381	l .
23048 534 23049 534		23182	1	837	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
ł	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		<u> </u>			744	LIE + GODGERD DOLL + DEDENIO
23050	53418	A	23184	3	/44	HEASCRSERRRQMADFDTYDD
1						RAYSSFGGGRGSRGS\AGGHGS
ľ						RSQKELPTEPPYTAYVGNLPF\N
						TVQGDIDAIFKDLSIRSVR\LVR
						DKVTGKFK\GFCYVEFDEV\DSL
			1			K\EALTYD\GALLGDRSL\RVDF
l						AKGRKTKIRVAFGF\RKGGPDD
	1					R\GFRDDF\LGGRGGSRPGD\RR
						TGPPMGSR\FRD\GPPLRG\SN\M
1			1			DFREPTEEERAQRPRIQ\LKPRT
						VWRPLNQVANPNSAIFGGARPR
ļ						EEVVQKEQE
23051	53419	С	23185	1	1038	
23052	53420	Α	23186	1	1486	MGRNQSRKAENSKHESTYSPP
		1				KDHSSSQAMEQSWTENGFEKL
						GFRRKEALYYLDLTASCQSQEL
	ľ	1				FQLYAGMSVVGTSMPVQAVCP
						YCGNRIITVTTFVPGALTWLLC
		ĺ			İ	TTLFLFGYVLGCCFLAFCIRSLM
	ľ	l				DVKHSCPVCQRELFYYHRLRQ
	ŀ					MELASSARTVVAPSRPCAGREG
	İ				ŀ	RGDANGKNSSPTAGSAMSSEPP
1	ŀ					PPPQPPTHQA\GVGLLDTPRSRE
			l .	l .		RSPSP\LRGNVVP\SPLPTRRTRT
			1			FS\ATVR\ASQGPVYKGV\CKCF
		ļ	l			CRSKG/HIGFNYPPAGWAGPDIF
	İ	l				PGTSLDVGRGSY\VPSGKGDEV
	ŀ	1				TLLKCAPIPTPKNE\KLAGPWEV
		1				VHHSPWHPGT\KHET\WSG\HVI
		1			i	SFLGGLVESTPLSLCFVGDFAG
		1				RROOTLEDDILPHETGLORGNG
		1				PSQVSPGGKGYGGAGVGCGVF
	İ	1			1	PAISTAYGPLOOPLHHLKSIKSH
		1	ŀ			LKRRGAHWWLSGGSNPIPGSGS
		1				RVGISSQLAQTHGLNPQNPSSS
	i	1				
22052	62421	<u> </u>	23187	2	617	WSWRPQIPRSGPLAA
23053	53421	A_		1	993	
23054	53422	A	23188	779	1210	A A A COST A CRIDED CA DISTRICTURA
23055	53423	Α	23189	1/19	1210	AAAGSEAGRRRGADPTRGPHV
				1		LPGSRQRRP/PIFPSLDLSGFFLF
	1			İ		CSEFCPEIKSTNPGISIGDVAKE
	1					KKKLGEMWNSLNDSEKQPYFT
						KAAKLKEKYEKDVADYKSKG
	1					KFDGAKGPAKVARKKVEEEDE
		<u> </u>				QEEEEEEEEEEDE

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nuclcotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23056	53424	A	23190	116	522	KAPYSTNSLSLPPFPILFMGSGY
23030	33424	ľ`	25170	1		RKGPTFQEPKPLPGAAARGPGC
		l				DPTQPPRHSAPRDHGPP\SPRCS
		l				PRPARPRASPAWLPPRAPTPPSP
		l	Į.			PAPAAAPRSCRGHARASSGRSG
		1	l			TTATPRRRRGGLWNQSPMDIK
		ı				G
23057	53425	A	23191	1	152	
23058	53426	A	23192	1	676	MWSAGRGGAAWPVLLGLLLA
22000	33 120	ľ.	25152	l'		LLVPGGGAAKTGAELVTCGSV
						LKLLNTHHRVRLHS\HDI\TYGS
		1				GSGOO\SVTGVEASDDANSYW
		1				RIRGGSEGGCPRGSPVRCGOAV
Į.	i	l		l		RLTHVLTGKNLHTHHFPSPLSN
1	l					NOEV\SAFGE\EGEGDDPGTLW
						TVRCSGTA/HWEREAAVRFQH
	İ	1	İ			VGTF\VFLSVTGEQYGSPIRGOH
						EVHGMPQCQ\QHNTWKAMEGI
						FIKPSVEPSAGHDEL
23059	53427	Α	23193	1	1044	
23060	53428	Α	23194	1	1083	MTATLAAAADIATMVSGSSGL
			i			AAARLLSR\SFLLPHNGIRHCAY
		l				TA/SLRQHLYVDKNTKIICQGFT
ł		i				GKQG/TPFHSQQALEYGTKLVG
		1				GTTPGKGGQTHLGLPVFNTVK
		1				EAKEQTGATG\SVIYVPPPFAAA
ĺ			1			AINEAIE\AEIPLVVC\YPRGIPQP
		1				GLGHSSSHKTCGPGKDKAYWG
	l	1	ľ			PNCPGV\INPGECKIGIMPG/HIF
	i				i	HKKGRIGIVSTSGTLTYEAGH/Q
1		l			1	TTQKLGLGQSLCVGIGGDP\FN
						GTDFID\CLEILFERFLPQEGIILI
					[G\EIGGNA\EENAAEILEPLYSGS
		L				KF/ARPVVS\FIAGLTAPPGRRN
					1	GVMPGAIISGREKVGAKRERSS
l	1		1			ALQSAGVVV\SMSPCTAGEPRF
l		1	1		1	YKEFEKREDRL

NO. of peptide sequence load on USSN ocodo for peptide sequence of peptide sequence of peptide sequence of peptide sequence of peptide sequence of peptide of pepti	SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
23061 53429 A 23195 1173 2110 FFTELEETILKFIWNQKRACISK		of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
ARVNTNOKSGGITLPOFKLYNE		- Interest					
AIVAKITWAWYSSRIHOQEKR REKK*KKDRSVPETLKKKQR NFTELNIQCLSKKFAQKMQIA RRKLIYEKVKIICHKEYRKMYR TEIRMARMATKAGNTYVPAEP KLAFVIRIGINGVSPKVRKVLQ LFLLYQIFINGAFVKLMASINM LRIVEPYIA/WRYPNLKSVNELI YKCGYGKINKK*IALTDNALTA RSLGKYGIICREDLHEITYTOK CFKEANNFLWPF/KLSSPRGGM KKKTHIFVEGGDAGKREVQIN RLIRRMN RLIRRMN RIFER RKKRAFAELKIKELRKKEVPAVPETLK KKRRNFAELKIKELRKKEVPAVPETLK KKRRNFAELKIKELRKKEVPAVPETLK KKRRNFAELKIKELRKKEVPAVPETLK RKKRNFAELKIKELKKEVPAVPETLK RKKRNFAELKIKELRKKEVPAVPETLK RKKRNFAELKIKELKKEVKEVPAVPETLK RKKRNFAELKIKELKKEVKEVPATLK RKKRNFAELKIKELKEKKEVKEVPATLK RKKRNFAELKIKELKEKKEVKEVPATLK RKKRNFAELKIKELKKEKKEVKEVPATLK RKKRNFAELKIKELKKEKKEVKEVPATLK RKKRN	23061	53429	Α	23195	1173	2110	FFTELEETILKFIWNOKRACISK
AIVAKITWAWYSSRIHOQEKR REKK*KKDRSVPETLKKKQR NFTELNIQCLSKKFAQKMQIA RRKLIYEKVKIICHKEYRKMYR TEIRMARMATKAGNTYVPAEP KLAFVIRIGINGVSPKVRKVLQ LFLLYQIFINGAFVKLMASINM LRIVEPYIA/WRYPNLKSVNELI YKCGYGKINKK*IALTDNALTA RSLGKYGIICREDLHEITYTOK CFKEANNFLWPF/KLSSPRGGM KKKTHIFVEGGDAGKREVQIN RLIRRMN RLIRRMN RIFER RKKRAFAELKIKELRKKEVPAVPETLK KKRRNFAELKIKELRKKEVPAVPETLK KKRRNFAELKIKELRKKEVPAVPETLK KKRRNFAELKIKELRKKEVPAVPETLK RKKRNFAELKIKELKKEVPAVPETLK RKKRNFAELKIKELRKKEVPAVPETLK RKKRNFAELKIKELKKEVKEVPAVPETLK RKKRNFAELKIKELKKEVKEVPATLK RKKRNFAELKIKELKEKKEVKEVPATLK RKKRNFAELKIKELKEKKEVKEVPATLK RKKRNFAELKIKELKKEKKEVKEVPATLK RKKRNFAELKIKELKKEKKEVKEVPATLK RKKRN							
REEKKE*KKDPSVPETLK*KOR NFTELKIQCLSKKFAQKMLQTA RRKLIVEKVKIICHKEYTKMYR TEIRMARMATKAQNFYVPAEP KLAFVIRIGINGVSPKVRKVU, L LFLLYQIFNGAFVKLNKASINM LRIVEPYIA/WKYPNLKSVNELI YKCQYGKINKK*IALTDNALTA RSLGKYGIICREDLIHEIYTVGK CFKEANNFLWPF/KLSSPRGGM KKKTTIIFVEGGDAGKREVQIN RLIRRINN 23062 53430 A 23196 3 571 GTMEOVEEKKKEVPAVPETLK KKRRNFAELKIKRI.RKFAQK MLRKARKLIVEKATA KKRRNFAELKIKRI.RKFAQK MLRKARKLIVEKAFAQK MLRKARKLIVEKAFAQK MLRKARKLIVEKAFAQK MLRKARKLIVEKAFAQK PVVPARTQIWRFVIRINGSINGS EPQRFRKVLQLLRPFVIRINGSINGS EPQRFRKVLQLLRPFVIRINGSINGS GNQFDLHFPTEKRSHPGSQ QNQFDLHFPTEKRSHPGSQ QNQFDLHFPTEKRSHPGSQ QNQFDLHFPTEKRSHPGSQ QNGFDLHFPTEKRSHPGSQ QYPVKDKKLLEVCTGGSPSPW ULMGOSSPSGIFGAFQRGLLR GTYNKYINVKKGSDLGGFTM LACYVLFSYSFSYKHLKHER! LRKYH LRKYH LRKYH LRKYH LRKYH LRKYH LRKYH LRGGRVLWM QACIMREYKLVVLGSGGVGKS ALTVQFVQGIFVEKYPPTIEDS YRKQVEVDAQQCMLELLDTAG TEQFTAMRDLYMKNOQGFALV VSTTAQSTFNIOLQDLREQURV KDTIDVPMILVGNKCDLEDER VVGKEQQQNLARQWNNCAFL ESSAKSKIVNEIFYDLVRQINR KTYPVPGKAPQKSPSCQLL ESSAKSKIVNNEIFYDLVRQINR KTYPVPGKAPQKSPSCQLL ESSAKSKIVNNEIFYDLVRQINR KTYPVPGKAPQKSPSCQLL SPHGEPLLQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCRREMWGWSLH TESPLCQGCREMWGWSLH TESPLCQGCREMWGWSLH TESPLCQGCREMWGWSLH TESPLCQGCREMWGWSLH TESPLCQGCREMWGWSLH TESPLCQGCREMWGWSLH TESPLCQGCREMWGWSLH TESPLCQGCREMWGWSLH TESPLCQGCREMWGWSLH TESPLCQGCREMWGWSLH TESPLCQGCREMWGWSLH TESPL							
NFTELKIQCLSKKFAQKMLQTA RRKLIYEKVKHCHKEYRKMYR RRKLIYEKVKHCHKEYRKMYR TEIRMARMATKAGNIYYPAEP KLAEVIRIRGINGVSPKVRKVLC LFLLYQIFINGAFVKLNKASINM LRIVEPYIAWRYPNLKSVMELI YKCQYGKINKK*IALTDNALTA RSLGKYGIICREDLIHEIYTYGK CFKEANNFLWPF/KLSSPRGGM KKKTHIFVEGGDAGKREVQIN RJIRRMN GTMEGVEKKKEVPAVPETKA KKRRNFAELKIKRLRKKFAQK MM.RKARRKLIYEKAKHYHKEY RQMYRIJERMAJMARKAGN FIVYPARTQIWREVIRINGRINGS EPQRFRKVQLQLRPFVLIFNGT FPVKLQQGFRINMLRDCRSHIL AWGYPNLKSVHELIYKJRGYG QNQPLHFPTEKRSHPGSQ QNQPLHFPTEKRSHPGSQ QYPVEDKKLLEVGTGGSPFSW ULMRGPSSPSGIFGVAFQGLR GTYYNKYINNYKKGSDLGGFTMV LACVYLFSYSPSYHILKHERE LRKYH LRKYH LRKYH LRKYH LRKYH LRYH						Į.	
RRKLIPEKVKHCHKEYRKMYR TEIRMARMATKAGNFYVPAEP KLAFVIRIRGINGVSFKVRKVLQ LFLLYQIFINGAFYKLNKASINM LRIVEPYIJAWRYPILKSVMELI YKCGYGKINKK*IALTDNALTA RSLGKYGIICREDLIHEIYTVGK CFKEANNFLWPFYLSVSPRGGM KKKTTIIFVEGGDAGKREVQIN RLIRRMN GTMEGVEEKKKEVPAVPETLK KKRRNFAELKIKKLRKKFAQK MKRKARKLIPKKAFAQK MKRKARKLIPKKAFAQK MKRKARKLIPKKAFAQK RQMYRITEIRMANMARKAGN FYVPARTQIWRFVIRIRGINGS EPQRFRKVLQLLRPFVKIFNG/T PFVKLQQGFRNMLRDCRSHIL AWGIYPNLLSVSPELIYKGRGY QNQFDLIFFIEKRSHPGSQ QYPVKDKLLLEVGTGGSFPSW ULMGOPSSPSGIFGAFQRGLLR GTYNKYINVKKGDLGGFTW LACYVLFSYSFSYKHLKHER LRKYH LACYVLFSYSFSYKHLKHER LRKYH LRYH LACYVLFSYSFSYKHLKHER LRKYH LRYH						1	,
TEIRMARMATKAGNIFYYPAEP KLAFVIRIRGINGVSPKVRKVI.Q LFLLYQIFNGAFVKLINKASINM LRIVEPYIAWRYPNILKSVNELI YKCGYGKINKK-MAIDNALTA RSLGKYGIIGREDLIHEIYTVGK CFKEANNFLWPFI/KJSSPRGGM KKKTTHEVEGGDAGKREVQIN RLIRRINN GTMEOVEEKKEVPAVPETI.K KKRRNFAELKIKRI.RKKFAQK MLRKARRKLIYEKAKHYHKEY RQMYRIVERMAIRMARKAGN FIVYPARTQIWRFVIRINGGINGS EPORFRKVJQLIAPPVIRINGGINGS EPORFRKVJQLIAPPVIRINGGINGS EPORFRKVJQLIAPPVIRINGGINGS ONQPEDLHFPTEKRSHPGSQ ONQPEDLHFPTEKRSHPGSQ ONQPEDLHFPTEKRSHPGSQ OYPVKDKKLIEVCYTGGSFPSW ULMIGDSSPSGIFGAFQRGLIR GTYNKYINVKGSDLGGFTM LRKYH							
LRIVEPYIA/WKYPNLKSVNELI							
LFLLYQIFINGAFVKLNKASINM	İ						KLAFVIRIRGINGVSPKVRKVLO
LRIVEPYIA/WKYPNLKSVNELI	ì						
RSLGKYGIIGREDLIHEIYTVGK							
RSLGKYGIIGREDLIHEIYTVGK							YKCGYGKINKK*IALTDNALTA
	1		1		i		CFKEANNFLWPF/K/LSSPRGGM
RLIRRNN							
						1	
	23062	53430	A	23196	3	571	GTMEGVEEKKKEVPAVPETLK
M.R.K.A.R.K.LIYEKA.K.H.Y.H.K.EY RQMYRT\EIRMA\RMARKI.AGN FIYVPARTQIMER\GYIRWARMARKI.AGN FIYVPARTQIMER\GYIRWARMARKI.AGN FIYVPARTQIMER\GYIRWARMARKI.AGN FIYVPARTQIMER\GYIRWARMARKI.AGN FIYVPARTQIMER\GYIRWARGA FIYVPARTQIMER\GYIRWARGA\GYIRWA			-				
FVYPARTQIWREVIRURGINGS EPQRFRKVLQLLRPFVKIFNG/T					i		
FVYPARTQIWREVIRURGINGS EPQRFRKVLQLLRPFVKIFNG/T							
PEQRERKVIQLLERPFKIENG/T PFVKLQQGFRINMLRDCRSHIL AWGIYPNLKSVNELIVKIRGVG QNQFDLHFPTEKRSHFGSQ QNQFDLHFPTEKRSHFGSQ QNQFDLHFPTEKRSHFGSQ QNQFDLHFPTEKRSHFGSQ QYPVKDKKLLEVQTGGSFPSW ULMKGPSSPSGIFGAFQRGLLR GTYNKYINNVKGSDLGGFTMV LACYVLFSYSFSYKHLKHGER LRKYH LRKYH PRVRLRC*MVDEYCGCEWGK* LNGGRVLWM LNGGRVLWM QACIMEPYKLVVLGSGGVGKS ALTVQFVQGIFVEKYDPTIEDS YRKQVEVDAQQCMLEILDTAG TEQFTAMRDLYMKNQGFALV YSTTAQSTFNDLQDLREQLIRV KDTDDYPMILVGNKCDLEDER VVGKEQGQNLARQWNNCAFL ESSAKSKIVNEIFYDLVRQINR KTYPYPGKAPQKSPSQCLL SPHGEPLLGQCRREMWGWSLH LS3066 53434 A 23200 461 1422 SPHGEPLLGQCRREMWGWSLH LS3067 53435 A 23201 232 331 TPPL*KWLRGVMAVTVYTPQTG TEPL-SWLRGVMAVTVYTPQTG TEPL-SWLRGVMAVTVYTPQTG TEPL-SWLRGVMAVTVYTPQTG TEPL-SWLRGVMAVTVYTPQTG TEPL-SWLRGVMAVTVYTPQTG TEPL-SWLRGVMAVTVYTPQTG TEPL-SWLRGVMAVTVYTPQTG TEPL-SWLRGVMAVTVYTPQTG TEPL-SWLRGVMAVTVYTPQTG TEPL-SWLRGVMAVTVYTPQTG TEPL-SWLRGVMAVTVYTPQTG TEPL-SWLRGVMAVTVYTPQTG TEPL-SWLRGVMAVTVYTPQTG TEPL-SWLRGVMAVTVYTPQTG TEPL-SWLRGVMAVTVYTPQTG TEPL-SWLRGVMAVTVYTPQTG TEPL-SWLRGVMAVTV							
PFVKLQQGFRFNMLRDCRSHIL AWGIYPNILKSVNELIYK\RGYG QNQFDLHFPTEKRSHFGSQ QNQFDLHFPTEKRSHFGSQ QNQFDLHFPTEKRSHFGSQ QNQFDLHFPTEKRSHFGSQ QYPVKDKKLLEVQTGGSFPSW ULMGDSSPSGIFGAFQRGLLR GTYNKYINVKKGSDLGGFTMV LACYVLFSYSFSYKHLKHERI LRKYH LRKYH LRKYH LRKYH LRKYH LRKYH LRKYH LRKYH LRKYH LRKYH LRKYH LRKYH LRKYH LRKYH LRKYH LRKYH LRKYH LRKYH LRYH							
QNQFDLHFTEKRSHPGSQ			1		1		
23063 33431					1		AWG\YPN\LKSVNELIYK\RGYG
QYPVKDKKLLEVQTGGSFPSW ULMKGPSSPSGIFGAFQRGLR GTYNKYNNVKKGSDLGGFTMV LACYVLFSYSFSYKHLKHŒR LRKYH LACYVLFSYSFSYKHLKHŒR LRKYH ENGERVLWM CACKMEPKLVVLGSGGVGKS LNGGRVLWM CACKMEPKLVVLGSGGVGKS ALTVQFVQGIFVEKYDPTIEDS YRKQVEVDAQQCMLEILDTAG TEQFTAMRDLYMKNQGFALV YSTTAQSTFNDLQDLREQLLRV KDTDDYPMILVGNKCDLEDER VVGKEQGQNLARQWNNCAFL ESSAKSKIVNEIFYDLVRQINR KTYPVPGKAPQKSPSCQLL SPHGEPLLGQCRREMWGWSLH CACKMEPKL CACKM			1				QNQFDLHFPTEKRSHPGSQ
NILMRGPSSPSGIFGAFQRGLLR GTYNKYRNVKKGSDLGGFTMV	23063	53431	Α	23197	2	340	PRVRPRVRDSKMASVGEVSGP
GTYNKYRNVKKGSDLGGFTMV			1				QYPVKDKKLL\EVQTGGSFPSW
LACYVLFSYSFSYKHLKfier			ļ.				\ILMRGPSSPSGIFG\AFQRGLLR
LRKYH			į .		1		GTYNKYI\NVKKGSDLGGFTMV
23064 53432			1				L\ACY\VLFSYSFSYKHLKH\ER\
LNGGRVLWM	1				l		
23065 53433 A 23199 123 693 QACIMREYKLVVLGSGGVGKS ALTVQEVQGIFVEKYDPTIES YRKQVEVDAQQGIFVEKYDPTIES YRKQVEVDAQQGMLEILDTAG TEQFTAMRDLYMKNGGGFALV YSITAQSTFNDLQDLREQILRV KDTDDVPMILVGNKCDLEDER VYGKEQGQNLARQWNNCAFL ESSAKSKIYVNEIFYDLVRQINR KTPVPGGKAPQKSPSCQLL SPHGEPLLGQCRREMWGWSLH TESPLGQWGSQILKLQNDLL 23067 53435 A 23201 232 331 TPL*KVLRGVMAVTVVTPQTG	23064	53432	Α	23198	2	48	PRVRLRC*MVDEYCGCEWGK*
ALTVQFVQGIFVEKYDPTIEDS YRKQVEVDAQQCMLEILDTAG TEQFTAMRDLYMKNQGGFALV YSITAQSTFN0LQDLREQILRV KDTDDYPMILVONKCOLEDER VVGKEQGQNLARQWNNCAFL ESSAKSKIYVNEIFYDLVRQINR KTYPYPGKAPQKSPSCQLL 23066 53434 A 23200 461 1422 SPHGEPLLGQCREBMWGWSLH TESPLGQWGSQILKLQNDLL 23067 53435 A 23201 232 331 TPL*KWLRGVMAVTVYTPQTG							
YRKQVEVDAQQCMLEILDTAG TEQFTAMRDLYMKNQQGFALV YSTTAQSTFNDLQDLEEQLIRV KDTDDVPMILVGNKCDLEDER VVGKEQQQNLARQWNNCAFL ESSAKSKIVYNEIFYDLVRQINR KTPIVPGIKAPQKSPSCQLL Z3066 53434 A 23200 461 1422 SPHGEPLLGQCRREMWGWSLH TESPLCIQWGSQILKLQNDLL Z3067 53435 A 23201 232 331 TPI.*KWLRGVMAVTVYTPQTG	23065	53433	Α	23199	123	693	
TEQFTAMRDLYMKNGGGFALV YSITAQSTFNDLQDLREQILRV KDTDDVPMILVGNKCDLEDER VVGKEQGQNLARQWNNCAFL ESSAKSKIYVNEIFYDLVRQINR KTPVPGKAPQKSPSCQLL			1	ł	İ		ALTVQFVQGIFVEKYDPTIEDS
VSITAQSTFNIDLQDLREQUIRV KDTDDVPMILVGNKCDLEDER VVGKEQGQNLARQWNNCAFL ESSAKSKIYVNEIFYDLVRQINR KTPVPGKAPQKSPSQLL 23066 53434 A 23200 461 1422 SPHGEPLLGQCREBMWGWSLH TIESPLCQWGSQILKLQNDLL 23067 53435 A 23201 232 331 TPL*KWLRGVMAVTVYTPQTG	1		1				
RDTDDVPMILVGNKCDLEDER VVGKEQQNLARQWNNCAFL ESSAKSKIYVNEIFYDLVRQINR KTPIVPGIKAPQKSPSCQLL 23066 53434 A 23200 461 1422 SPHGEPLLGQCRREMWGWSLH TESPLCIQWGSQILKLQNDLL 23067 53435 A 23201 232 331 TPL*KWLRGVMAVTVYTPQTG	1		1		ł		TEQFTAMRDLYMKNGQGFALV
VVGKEQGNLARQWNNCAFL ESSAKSKIYVNEIFYDLVRQINR KTPVPG0KAPQKSPSCQLL 23066 53434 A 23200 461 1422 SPHGEPLLGQCRREMWGWSLH TESPLGQWGSQILKLQNDLL 23067 53435 A 23201 232 331 TPL*KWLRGVMAVTVYTPGTG			1		1		
ESSAKSKIYVNEIFYDLVRQINR KTPVPGKAPQKSPSCQLL	1					1	
KTP\VPG\KAPQKSPSCQLL 23066 53434 A 23200 461 1422 SPHGEPLLGQCREMWGWSLH TESPLG\03000 TESPLG\	1			l	1		VVGKEQGQNLARQWNNCAFL
23066 53434 A 23200 461 I422 SPHGEPLIGOCREMWGWSLH 23067 53435 A 23201 232 331 TPL*KWLRGVMAVTVYTPQTG						l	
TESPLG\QWGSQILKLQNDLL 23067 53435 A 23201 232 331 TPL*KWLRGVMAVTVVTPQTG			L				
23067 53435 A 23201 232 331 TPL*KWLRGVMAVTVVTPQTG	23066	53434	A	23200	461	1422	
			L				
I I I I I I I I I I I I I I I I I I I	23067	53435	Α	23201	232	331	TPL*KWLRGVMAVTVVTPQTG
IIAII DI MILWOSTOTI KETA			L				IIAIPDPHHEWGSTGYPRLPA

SEO ID	SEO ID NO-	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23068	53436	A	23202	5	1223	I GLDSGPESGLRAQPGLLAGKRK
23008	133430	^	23202	ľ	1223	MSVLRRMMRVSNRSLLAFIFFF
	1				1	SLSSSCLYFIYVAPGIANTYLFM
	1	1				VOARGIMLRENVKTIGHMIRLY
	į.	l				TNKNSTLNGTDYPEGNNSSDYL
		1				VOTTTYLPENFTYSPYLPCPEKL
		1				PYMRGFLNVNVSEVSFDEIHOL
		1				FSKDLDIEPGGHWRPKDCKPR
		1				WKVAVLIPFRNRHEHLPIFFLHL
		1				IPMLQKQRLEFAFYVVEQTGTQ
				1	1	PFNRAMLFNVGFKEAMKDSV
						WDCVIFHDVDHLPENDRNYYG
1		ŀ				CGEMPRHFAAKLDKYMYILPY
						KEFFGGVSGLTVEQFRKINGFP NAFWG\WGGEDDDLW\NRV\H
		1				YAGY\NV\TRPEGDLGKYKSIPH
					ŀ	HHRGEVQFLGRYKLLRYSKER
		1				QYIDGLNNLIYRPKILVDRLYT
		1				NISVNLMPELAPIEDY
23069	53437	A	23203	78	495	CTFEKAPPFOLGRGRMVPAKK
23009	33437	^	23203	/°	493	GGEKKTGR\AINEVVTREYTINI
		1				HKRIHGVGFKKRAPWALKEIR
		1				KFAMKEMGTPDVRIDTRLNKA
		1	ł			VWAKGIRNVLYRIHVRLSRKR
		1				NEDEDSPNKLYTLVTYVTVTTF
		ı			ì	KNLOTVNVDEN
23070	53438	A	23204	138	360	KNOKROLSGGHCADOTSLVLA
23070	33436	<u> </u> ^	23204	138	500	RLAPLF\PTTMSDKPDMAEIEKF
	1	ı				DKSKLKKTETQEKNPLPSKETT
		1				EQEKQAGKS
23071	53439	A	23205	145	728	KRQPTSAMKDPSRSSTSPSIINE
25071	33437	ľ.	20200	145	/20	DVIINGHSHEDDNPFA\EYMWD
1	ŀ					GKMEGRNFNR\OIGRRELWGR
1		1				KD\FLNRCFPKE\MLGEEGKEG
	1					ALNWVLFPSFEILPHOTMGPNP
	İ	l				RTSFNDPCLSSDGSFSWEDLVV
	į.		i	ļ		KS\NLNPNAKEFVPGGEVRKYL
	l					SRRGPLLVDVAQFPHCEGSIRR
		1		i		LNCKSSLGHCVTLYAIAQSFC
23072	53440	A	23206	530	707	Quic
23073	53441	A	23207	1	515	MDFSKVPPHAAFRAQPPPALT
[ľ	l	ľ		WGLAKGSGRGAGEGGERPDRR
						R/PAARTGTPAWRPAGRRRLPP
	1	l	1	I		RSPPPAPVPVRPALPHPVRPGKE
	1	l	1	I		TGKTRRRRHYSPGRPLRGAAA
	1	l		I		AAAAAAVAOTASPSWAEGGRS
1	1	1		I		GAGGSAESPAFSSPALPCPEGLL
				1		CPSAVPVPSKQSVSEVLAIF
23074	53442	A	23208	3	713	C.C TIGINGO TOD TOAIL
23075	53443	Ā	23209	1	704	
23073	1555	٠.	20207	ı	/ / .	

	01/0/306/					PC1/US01/08631
SEQ ID	SEQ ID NO:					Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence	ı	09/540,217	sequence	or peptide sequence	deletion, (-possible nucleotide insertion)
				o-quemee		
23076	53444	A	23210	2	1631	GLPRRPTRPPTRSGVSPPWPGPF
						SGVLFPLLLHCTGSLSQAVLTQ
1		l			1	PSSLSASPG\ASVKLTCTLRSDIN
		l				VGSPFIYWYQQKPGSPPQFLLR
		l				YKSDSDNQQGSGVPSRFSGSKD
		Į			i	ASANAGILLISGLQSE\VRDDYY
		1				CQAWDSSTAHSDTGRCGIVGRI
		1				PVRQCWPLLLLGLAMVAHGLL
						RPMVAPQSGDPDPGASVGSSRS
		İ		1		SLRSLWGRLLLQPSPQRADPRC
						WPRGFWSEPQSLCYVFGTGTK
						VTVLGOPKANPTVTLFPPSSEEL
						OANKATLVCLISDFYPGAVTVA
						WKADGSPVKAGVETTKPSKOS
1						NNKYAASSYLSLTPEQWKSHR
						SYSCOLOWGLGLGASQGGFLY
						EPVSQCVVFGGGTKLTVLGQP
						KAAPSVTLFPPSSEELQANKAT
						LVCLISDFYPGAVTVAWKADSS
1						PVKAGVETTTPSKQSNNKYAA
1						SSYLSLTPEQWKSHRSYSCQKP
						NHASHGTSRMPSVSAPVSGLKS
		1				CWSGTGDRSALGQMPGPCHGI
1						PDPMQQASDSGQPKAAPLGSL
22000	53445	ļ	22211	1	959	CSKPSSEELSGQQGPHWCVS
23077	33443	Α	23211	[1	959	VFLLLYLVAMPDPAKSAPAPK
1		ŀ				KGSKKAVTKAQKKDGKKRKR
						SRKESYSVYVYKVLKQVHPDT
1						GISS\KAMGIMNSFVNDIFERI\A
1		ļ			1	GEASRLAHYNKRSTITSREIQTA
]						VRLLLPGELAKHAVFIENLLNY
1						RKYLVKSSEKDRNYPCPLPGLK
1						SKVDICTPYCLLLLPQPRHYPG
			ĺ			KDTKEETEDFPNSPPPINREEDK
						RYTTVMGPCLRQAALEGELLA
						CPVMEDQQANWVHEPITFNTF
						KEIRKSIRENGPASPFTRGLIEVI
1				l		ADSSDLVASTQKLTQPKETPSA
	1					PYNDISDLTNQDSWFIGFPPIWN
						AGETDLSNNKTT
23078	53446	A	23212	3	463	
23079	53447	A	23213	74	464	
23080	53448	Α	23214	41	490	
23081	53449	A	23215	2	369	PDPAANAPAAAKPAAAKALPV
			1	l		FLLLLSGGAGPAHPMRAPGC*G
1	l					ELGYTEQLFQFLQATAQEEGIF
1	l	İ				EYTDPKLAISAVWTFRDLGVLQ
1	l	l				QTPSPAGPRLHLSPTFASLDNQE
						KLEQFIRQFICS
23082	53450	A	23216	124	301	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nuclcotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide scquence	delction, \=possible nucleotide insertion)
				sequence		
23083	53451	A	23217	3	2426	PGVLSTATFRESGLSPCSKLKHP
						GPRGRLOYSPPTAGOGRLENG
						AGHPOSTGPVATLSVAEYGTO
		l				GGASGPARIPIGSRGCLLAGRA
						EASLWSSGFGMKLEAVTPFLG
			l			KYRPFVGRCCQTCTPKSWESLF
						HRSITDLGFCNVILVKEENTRLF
			ŀ			SWALLRVLKLLFLNVQLHKGQ
						MKMVQKAAQQVGQVGTSLWL
						SPEOGLPLVLLSTHKTLLDGILL
			ŀ			PFMLLSOGLGVLRVAWDSRAC
			l			SPALRALLRKLGGLFLPPEASLS
						LDSSEGLLARAVVOAVIEOLLV
		ļ	ŀ			SGOPLLIFLEEPPGALGPRLSAL
		1				GOAWVGFVVQAVOVGIVPDAL
		l				LVPVAVTYDLVPDAPCDIDHAS
		l				APLGLWTGALAVLRSLWSRWG
		i i				CSHRICSRVHLAOPFSLOEYIVS
		l l				ARSCWGGRQTLEQLLQPIVLGQ
		1		İ		CTAVPDTEKEOEWTPITGPLLA
		l l				LKEEDOLLVRRLSCHVLSASVG
		l				SSAVMSTAIMATLLLFKHOKG
		l				VFLSQLLGEFSWLTEEILVRGFD
		l				VGFSGOLRSLLOHSLSLLRAHV
		l				ALLRIROGDLLVVPQPGPGLTH
		l				LAOLSAELLPVFLSEAVGACAV
1		l				RGLLAGRVPPOGPWELOGILLL
		l				SQNELYRQILLLMHLLPQDLLL
1		l				LKPCOSSYCYCOEVLDRLIOCG
		l				LLVAEETPGSRPACDTGRORLS
		l				RKLLWKPSGDFTDSDSDDFGE
		l				ADGRYFRLSQQSHCPDFFLFLC
23084	53452	A	23218	169	955	ASSQLPCSMAQCCATWAPLKM
		l				RWLCSTAWSTSSLSLPPSWSSL
		l				QASQPSCCHATSLAPPCAGQCL
		l	i			ARAWPVLSFLCTCALGLLASIA
		l				MTFATOGKALLAACTFGSSELL
		l				ALAPYCPFDPTRIYSSSLCLWGI
		l				ALRCSAWAENVFAVRCAOLTH
		1	l	l		QLLELRP\WWGKKQPTHMMRE
			l	l		NPELVEG\RDLLSCT\SSE\PLTL
		l		1		CRDDVLPRP\DGQLRTLQATLL
				l		CDPGPGFLGAGLRGLKWTLGG
		l	l	l		P\SGAFQPSSPTTPAHLHLKIETL
		l				F
23085	53453	A	23219	h	344	
25005	100.00	٠.	22217	L	2	

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence	l	02/340,217	sequence	or peptide sequence	deletion, (-possible intereorde insertion)
		1				
23086	53454	Α	23220	214	1355	FCSSWLRSLADSSLSWKMFLVC
	ł	1				LTGGIASGKSSVIQVFQQLGCA
	ŀ	1				VIDVDVTARHVVQPGYPAHRRI
	ŀ	l				VEVFGTEVLLENGDINRKVLGD
	ľ	l				LIFNQPDRRES\LNAITHPEIRKE
	Į.	l		ì		MMKETFKYFLRGYRYVILDIPL
		l		i .		LFETKKLLKYMKHTVVVYCDR
	1	1				DTQLARLMRRNSLNRKDAEAR
	1	l				INAQLSLTDKARMARHVLDNS
	1	l		ľ		GEWSVTKRQVILLHTELERSLE
		l				YLPLRFGVLTGLAAIASLLYLLT
		l				HYLLPYAYTASAIPPSGAASRG
		l		1		PGARCSAIRKRAPARPTSAVSPP
	ŀ	l		l .		GAPRSPARGLRAIDRAAGTRNL
		l				NATLPPPGPPAPPSPPSPPSSITL
	l .	l				HVFHQLQPDITRGREGTDFQWE
				l		KFPGRVLVADSRRALRCLAPAP
				·		TWLV
23087	53455	Λ	23221	2	1231	TPLFLRKGTADGSARTTVIHLG
		ł	:	l		FTKGKQRCVETQAAAVAGDVG
	ì	l	1	l		RTAAGGRVGKESGVISSDDRPG
	l	1	İ			LPVGQSQPEGGEQEATDTVSVH
		l	1	l		ATTSQAWPGGADSKELSSKAE
		1		ŀ		ERETGKDQQTELPLCLPLRDFP
					l .	DSAPVMSSDSGHATESLHLASS
				l		DSKPPMPTPNEGPGSTVGHAGP
				l	l	PRTVKADLCTEPQLALRGLQVS
					1	LPRLHLVGVIYAEHQASGLEGC
				l		GHAGSCSPDGSPRAPGATLCLS
				l	1	KPLSFRTKRCLPCWLPASAPVGI
				i		PGAPVHIPGAPVCIPGAPACISN
				l .		APVHIPGAPVGIPGAPVCIPGAA
		1	İ	l		VCVPNTPMCIPSAPMCIPNTPVC
		1	1	l	1	IPNTPVCIPNTPVCIPGAAVCILG
		l	I	l		A/PRAHPWCSSCVHPGAPECYL
		l		ŀ		GCGLTQADRLEPAVLDAGLQE
		<u></u>				PLFALCGQALPS
23088	53456	Α	23222	2	408	
23089	53457	A	23223	62	251	

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence	İ	
		<u> </u>				
23090	53458	A	23224	2111	3003	TQKAICMCLGLCKKIHHLLSLQ
1	l					CLISQIKMIMGAGWQ/SSGFTDE
1	l	1				KETSKIKADPEKLKKLQESLLE
1	l					EDSEEEGDLCRICQIAGGSPSNP
1	l	1				LLEPCGCVGSLQFVHQECLKK
		l			Į.	WLKVKITSGADLGAVKTCEMC
l					i e	KQGLLVDLGDFNMIEFYQKHQ
		1				QSQIKTEATVVEHKDIYNVTET
						QLVPEARSVLLVGAGQMLLTG
i						ANRALNSDYSSMRPKVKAQNE
					İ	LMNSGLYLVLLLHLYEQRFAEL
		l				MRLNHNOVERERVRVTRVLGG
1		1				LSPNVSCGVVSGVGSFQWVRG
		1				LTDLKNEASDLHAQIPAE
23091	53459	A	23225	ī	594	MIAAFNPGKVDTVTTNGPFIDL
		1				NYMVYMFHYDPTHGKFHDVP
1		l				MFVMGMNHKHENKLTVISNAS
		1	i			CTTNCLDPLAKVIHDNFGIREG
1		1				LMTTVHSIIATQKTVDGPSRKL
					Į.	WHDSHGAFONTIPASTGAAK\I
l		l				KKVVRQA/SQGPLKGILGYTEH
		1				OVVPLEFKSNTRPSTFEAGAGI
				i		AVNDHFVKLISWYGNEFAYSN
						KVIKPM
23092	53460	Α	23226	3	232	
23093	53461	Α	23227	1078	2221	QQTLQPAALPCPFPPPTHPPEVL
					1	PTAMNTTSSAAPPSLGVEFISLL
	l					AIILLSVALAVGLPGNSFVVWSI
l					ŀ	LKRMQKRSVTALMVLNLALAD
						LAVLLTAPFFLHFLAQGTWSFG
						LAGCRLCHYVCGVSMYASVLL
						ITAMSLDRSLAVARPFVSQKLR
						TKAMARRVLAGIWVLSFLLAT
i					ŀ	PVLAYRTVVPWKTNMSLCFPR
						YPSEGHRAFHLIFEAVTGFLLPF
						LAVVASYSDIGRRLQARRFRRS
		1				RRTGRLVVLIILTFAAFWLPYH
1		1		1		VVNLAEAGRALAGQAAGLGLV
1		1	l	I	ŀ	GKRLSLARNVLIALAFLSSSVNP
1		1	1	ŀ		
l		1		1		VL\YACAGGG\LLRSAGVGFF\A
				l		KLLEG\TGSEASSTP\RGGSLGQ
l			l			TV\KSGPAALEPGPSESLTASSP
		١.		6105	(212	LKL\NELN
23094	53462	A	23228	6125	6319	ASVVPVTSALAMCSPKSGIGDG
l	1	ı	l	1	l	FAVLRVEDVAQK***KAALGV
22225	52462	١	2222		1242	ASSPA WQQTREREHVQGNSPL
23095	53463	Α	23229	I	1347	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23096	53464	A	23230	1	1124	MRECISIHVGQAGVQIGNACWE LYCLEHGIQPDGQMPSDKTIGG
		1	İ			GDDSFNTFFSETGAGKHVPRAV
		1				FVDLEPTVVDEVRTGTYRQLFH
		1				PEOLITGKEDAASNYARGHYTI
1		1				GKEIVDLVLDRIRKL\FGGGTGS
1		l	l			GFASLLMERLSVDYSKKSKLEF
1		l	ŀ		ľ	AIYP\APQVSTAVVEPYNSILTT
						HTTLEHSDCAFMVDNEAIYDIC
		1				RRNLDIERPTYTNLNRLIGQIVS
1		1				SITASLRFDGALNVDLTEFQTN
						LVPYPRIHFPLATYAPVISAEKA
ł						YHEQLSVAEITNACFEPANQMV
		1				KCDPRHGKYMACCMLY\RGDV
				i .		\VPKDVNAAIATIKTKRTIQFVD
1						WCPTGFKV*LGDMEAFQQAAD
1						AQNNTGLEGPHPLGRLSLFHGL
23097	53465	Α	2323 I	I	2918	MDVYCAHESLRHTKSPEATGQ
		i				TSGQRPEALQAFWPAFGSSPVT
1		i .				GSSPVTEVTTKGRWSSSRQVL
İ		1				WDRIPLAKSRPAISNRQSQRTNI
		1				PWWESCPPILNLEREQPPRFAQP
		l				GTYASRWKALEEMEKQQREQ
		1				VDRNIREAKEKLEAEMEAARH
		1				EHQLMLMRQDPGLALVSCMQ
	İ				İ	QGQSMSRSPKKVFKHLYELNL
					Ì	TKLQSLGSRSRLRWKDSPFWGE
		į .	1		l	EAGGKRPPLVSLALHRSPSLLL
22000	60466	<u> </u>	22222	,	158	LFVPKHPSKGRPDCTVQRS
23098	53466	A	23232	1	414	LGGLKGTWGETKPAMAAAEEE
23099	33407	^	23233	'	414	DGGPEGPNRERGGAGATKTLL
		1				YTGNHLFSPOWLETRPEROECP
					1	VCKAGISREKVVGEKSLLTFSL
						HFLRLKTPOICRLPV*LFLPP*GF
						OPFGDTGGFHFSFGVGAFPFGF
		1				FNSPSY
23100	53468	A	23234	1	663	
23101	53469	A	23235	2	667	SWAGGGFWFVGACVCVFGGTE
		1		[_		GYVGRNKTGHGSSRRRRSGAP
		1				NVPNRERGGAGRPKTPFECNIC
		1	1			LETARELVLSVLANLYFWTISH
					1	QV\LE\TRPE\RQECP\VCKAGISR
		1			I	EKVVP\LYGRGSQEAPRIPRLKT
		1			1	PPPPPRGQR\PAPE\SRGGF\QPFG
	1	1			I	DTGGFHF\SFGVGAFPFGFFTTV
					1	F\NAHEPFRRGTGVDLG\QGHP
					1	A\SSW\QD\SLLLFLAIFFFF\WLL
23102	53470	A	23236	363	476	
23103	53471	В	23237	224	364	

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \"possible nucleotide insertion)
				sequence		
23104	53472	Α	23238	1	1324	MRQRRAELLSKDKRVRTDGNN
						KCSPSAIGKRKERGTNDWIEGD
		l				NNQESQGLNCSDTLNRDLGPN
i		l		i		TRGFLYAGLSGLDPDPSLPTPD
		1				MSSEVLEDNLDTLSLYSGKDSD
		1				SVKLLEEYADSESQASLQGRSR
						CWQHPHPAPLSYPPAASGSQLS
						LQGKFPSAKPPLLGGYTCTAQR
				l		TQWALTKRSLRATRESVSRQQ
						GVYGLVGRKRTRHHRGVLGLL
		1				GIPSARPERPGTGSEAGPTRLPG
						AAVARRRPARPVRGRLGDPAP
1						TAAQTSGPASLPPPPPPRSYGDP
						HONGYPDPDPAPSASPDLPPAA
					l	GDTRGSGGRRKKRRTGTARLN
						TPGRKA\PKQKAAAGRKGRAH
						OPAPRTPEPORRRPIRPPNOTOR
	i					LRHPHRRAGRGPPPGGAGGTA
						NPSTGANGGAATARRTAGART
						RGARGASERAGRHGSRPASWL
	İ					TATPPPGT
23105	53473	A	23239	1	433	PTRPSSPEETDGPWDPEGHPGG
25105	33473	ľ	23237	ľ	455	PLTDPPNAPVLALPPAIPGLSLS
		1				MADGSSDAAREPRPAPAPIRRR
						SSNYRAYATELODLCROLHAR
					i	VDKVDEERYDIEAK\VTKNITEI
						ADLTQKIFDLRGKFKRPTLRRV
						RISADAMMQALL
23106	53474	Α	23240	2	409	RISADAMINQAEE
23100	53475	Ā	23240	3	490	SSPEETVGPWDPEGHPGGPLTD
23107	33473	<u> </u> ^	23241	ľ	450	PPNAPVLALPPAIPGLSLIMADG
		1				SSDAVRAAG*GVAGTPRARAR
					1	EPRPAPAPIRRRSSNYRAYATEP
		1				HAKKKSKISASRKLOLKTLLLO
	1	Į.				IAKOELEREAEERRGEKGRALS
		İ	į			TRCQPLELAGLGFAELQDLCRQ
i		1			1	
	******	<u>. </u>	02242	1.40	046	LHARVDK
23108	53476	Α	23242	149	846	VSAWRNGSSDAAREPRPAPAPI
1		l		1	1	RRRSSNYRAYATEPHAKKKSKI
		1		I		S\ASRKLQLKTLLLQIAKQE\LE
		1			1	REAEERRGREGARSLAPRC\QP\
		l	1	1	1	LELAG\LGFADLPGLCCRQFHA
1		1				RV\DKVDEERYDIEAKVSKNITE
	1				1	IAN\LT\QKIFDLRSKFKRPTL\RR
	1	l	1	1	I	VRISADAMMQALLGARAKESL
	1				1	DLRAHLKQVKKEDPEKENREV
	1	1		1	1	GDWRKNIDALSGMEGRMKKW
1						
1						RELSLPAYCPCPEEGH
23109	53477	A	23243 23244	1 80	487	RELSLPA YCPCPEEGH

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, ^=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23111	53479	A	23245	427	1417	KTMWKVEFGVRFSITRLLEHQ PKQTMYIPLPK/PPLWPEEPKTE APKQTPRAPPKPKTSPRPRIPQT QPEETDQSTQEPFTTKIPRTTEL
}						AKTTQGLEMLSCLIHNPKMHK
						DSTNEQIAASLQACYYEQLLM
						YIRVTDVDSVGPGNGKEAVTA
ŀ						PQAPTLFWLNHKAQLAKDWA
						GKDVNITPTFATSKKENKPPNN
						LQICNVSKIHQRAKVAGKPEIK
						DKHIQKEMPLDAIEASKINLVKI
						FNEFLKAWKCRLSFEYRIPENQ
						GHKGSFYLLVGSSRTSWLLTEK
		1				IGSSKRVLRMPPMDILKTTQILL
		_				SFLWSINLTPRGKNNHLVAPAH
23112	53480	Α	23246	3	1011	
23113	53481	A	23247	2	334	PIQSFFLVLKKPPLRKGVYLSSK SLSRWFFKVQAN*RSQPQKAFS RLSCHC
23114	53482	١.	23248	1	642	RESCRE
23114	53482	A A	23248	854	1150	KEDTNKWKNIPCSWLGRINIVK
23113	33463	^	23249	0.54	1130	MAIMPKVNL*IOCHPHOATNDF
ì		l				SSQNWKKTTLKFIWEPKKEPAL
		l				PSQS*AKRAKLGGITLPDFQTIT
1		l				YKATVTKTA
23116	53484	A	23250	1754	2393	PSNRREFGESWPETQAPGALRP
23110	33404	ľ`	23230	1,754	12373	LLPAPGTPRARERPAGWPARPM
	ľ	1				EVYIPSFRYEESDLERGYTVFKI
ŀ		l				EVLMNGRKHFVEKRYSEFHAL
İ		1				HKKLKKCIKTPEIPSKHVRNWV
i		l				PKVLEQRRQGLETYLQ\AVILE
1		1				NEELPKLFLDFLNVRHLPSLPK
		l				AESCGSFDETESEESSKLSHQPV
		1				LLFLRDPYVLPAASDFPNVVIE
1		1				GVLHGIFYPHLQPR
23117	53485	A	23251	1	265	AGNRLRFQLELEFVQCLANPNY
		1				LN/WYPQCLHMLELLQYEHFR
		1				KELVNAQCAKFIDEQQILHWQ
1		l				HYSRKRMRLQQAL\AKQQQQN
		1				NTSGK
23118	53486	Α	23252	1	987	
23119	53487	Α	23253	1	491	RGFRNFARVSGLLLCQAGGVL
l l		1			l	VSSFVMAAAVAMETDDAGNR
		1				LRFQLELEFVQCYANHNTLIFLP
1		1			1	QR\GYF\KDKAFVNYLKYLLYW
1					1	\KDPEYAKYLKYPQCLHMLGA
1		1	l	1	1	APNMEHFRKEAGWNAQC\AKF
						IGWNRQVSTLGKHYS\RKRM\R
L			L			LQQALAEQQQQNNTSGK

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, /=possible nucleotide insertion)
23120	53488	A	23254	73	355	HEGOVHVLLASAHVDVAIVLG
23120	33400	1	23234	1,3	333	AGSMVEGEARPEOAAOHHOGI
		1				APLHLDAPGPSWTOSRSPRNRA
		1				NONHTCSEAT*SHPEOLPOTLW
		ı				ECHAPRSP
23121	53489	В	23255	1	441	
23122	53490	Ā	23256	528	698	HGQRFLHGPV/H/HHSHYDN/PH
						HSLHHHRCLHHCHLHHHRHHL
						HHHPLHCYHHHPGQNG
23123	53491	Α	23257	1	403	
23124	53492	Α	23258	509	4547	LYLTTSRREEHLKSLVVETQDL
						ASPVLRSVSICTKVEEAFRQAH
1						VIVVLDDSTNKEVFTLEDCLRS
						RVPLCRLYGYLIEKNAHESVRV
		1				IVGGRTFVNLKTVLLMRYAPRI
1		1	İ			AHNIIAVALGVEGEAKAILARK
						LKTAPSYIKDVIIWGNISGNNYV
1				٠.	•	DLRKTRVYRYESAIWGPLHYSR
					ľ	PVLNLIFDSEWVKREFVAILKN
						LTTTGRQFGGILAAHSIATTLKY
ł						WYHGSPPGEIVSLGILSEGQFGI
						PKGIVFSMPV
23125	53493	A	23259	56	290	
23126	53494	A	23260	468	993	VLLACFWPGSDFWPRSRRKTY
					1	GTAPQSCYSFYVALDCGGVSRL
l	İ					VIFVSWRNPQVAPTSAHQNRPS
f	ł				i	RNPVSRPPNTQRVARRKHYAL
ľ	1		l	l		ADGYTERRWTNAPCRAESSFPP
ŀ			l			NCPSAAPIPDSYDK*PSRATLFT
						SHLTVEAFPVWSYLYPGETPRL
ľ						HPRAHTRTVPAAIRYPDRQTPN
				i		GLRGANTMHSQMDTLNGDGR
ŀ						THLAGQRARSPRTARPQRQYRI
1						HTTSDLPGLVRKYASGCCLRIW
İ		1				YCCAWIERALSLQVGGGVFGD
23127	53495	-	23261	1	3114	PLSEGQAECDHGST
23127	53495	A	23262	3	401	TSSLSLSGKSGRYIVFLRRSVGI
23126	23490	<u>۱</u> ^	2.3202	2	401	OSPSAVATVRLLLAGSDRRFAA
1		l	1			GSAGCAVLSRAERS*EPGCCYR
1	l	İ				IRRSAFRLRAGSIGNRDKPVAW
	l	l	I			PDTGSGDNTEVYFRAHQAGILP
l		l				DTVLAGPAQRVAVSGGTPVDW
23129	53497	A	23263	2874	3062	DI VEAGI AQRVAVSOGIT VDW
23129	2347/	<u></u>	23203	2014	3002	L_,,

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23130	53498	A	23264	2766	3323	MPDLVEGREKSDKGAQPQPRT QSRSARTNGSGVRILHSLWRIIG IPLLLGYSLVCSRVLLACFWPG SDFWPRSRRKTYGTAPQSCYSF YVALDCGGVSRLVIFVSWRNP QVAPTSAHQNRPSRNPVSRPPN TQRVARRKHYALADGYTERR WTDAPCRAESSFPPNCPL*RQY RIHIVDASVCGP
23131	53499	Α	23265	968	1440	
23132	53500	A	23266	3	640	LSVVRSRIHMGVGVGCPVR YRNQDDHELQTHGNKIL.CGIV CDKGAQPQPRTQSRSARTNGSG VRILHSLWRIIGIPLLLGYSLVCS RVLLACFWGSDFWPRSRRKT YGTVPOSCYSFYVALDCGGVS PRRNPVSRPPNTQRVARRKHYA PADGYTERRWTIAPCRAESSFP PNCPSAAPIPDSYDK
23133	53501	A	23267		3140	MVYKMRYRSOHPYSIKEKOMK SEVLSVKEKIGYGMGDAASHII FDNVMLYMMFFYTDIFGIPAGF VGTMFLVARALDAISDPCMGL LADRTRSRWGKFRPWVLFGAL PGIVCVLAYSTPDLSNNGKMI YAAITYTLLTLLYTVVNIPYCA LGGVITNDPTQRISLQSWRFVL ATAGGMLSTVLMMPLVYLIGG DNXPLGFGGDLSPMFSTPEEIA RCPPYENDVHVGASSLAAGH KTLIPELVRSAEQHMGTR
23134	53502	A	23268	750	1340	ILHAPAPPESASASHEQFEWSD KGAQPQPRTQSRSARTNGSGVR ILHASLWRIGIPLLLGYSLVCSRV LLACFWPGSDFWPRSRRKTYG TAPQSCYSFVVALDCGGVSRLV IEVSWRNPQVAPTSAHONRPSR NPVSRPPNTQRVARKHYALA DGYTERRWTNAPCRAESSFPPN CPFAAPIPDSYDK-PFRIVS
23135	53503	A	23269	93	699	RASVQOKKLSTIDKGAOPOPR TOSSRARTNOSGVRILHSI, WRII GIPLLLGYSLVCSRVLLACFWP GSDFWBRSRKRTYGTAPOSCVS FVYALDCGGVSRKRSYLYPOET FRVAPTSAHONRPSKROTTRPP NTORVARRKHYALADGYTER SATNAPCRAESSFPPELPVPQVP NTGFIRQVTFRDCFVNTPQAAA SGFW

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23136	53504	Α	23270	1000	1530	QRYSDNPPNDKGAQPQPRTQSR
						SARTNGSGVRILHSLWRIIGIPL
	ł					LLGYSLVCSRVLLACFWPGSDF
						WPRSRRKTYGTAPQSCYSFYV
		l				ALDCGGVSRLVIFVSWRNPQV
1		l				APTSAHQNRPSRNPVSRPPNTQ
				i		RVARRKHYALADGYTERRWT
		l				NAPCRA/VEPFPPNCPWEAQYRI
23137	53505	Α	23271	242	711	LKQKRDKEDRSSRVSAVLLIK/
		l				HRRPLPLPIGLKACHCSCTAKC
l	1					LGGCESPRKGGQLLIFEGHVKI
ŀ		l				DLEIFTKSLWLTRQLIVKKETEK
1						AVKHASNICIMRTGLTFQSHKIE
		l				DGLFGLLVIATEAEWSYVCOST
l	ŀ	l				NPVDVYPLTAVPACCIISASDVP
23138	53506	A	23272	2	459	CVNSOIGRGHKITTYPFSPRKSG
		l				RKGMAOSOGWVKRYIKAFCK
	l	l				GFFVAVPVAVTFLDRVACVAR
l		l				VEGASMOPSLNPGGSQSSDVVL
ŀ						LNHWKVRNFEVHRGDIVSLVS
l		1	1			AKNPEOKIIKRVIALEGDIVRFP
	İ	l				*DFCMPMPHISCGPQSAGRNW
23139	53507	A	23273	1	474	
23140	53508	Α	23274	358	948	PLTPFSPRKSGRKGMAQSQGW
1		1				VKRYIKAFCKGFFVAVPVAVTL
	1		1			LDRVACVARVEGASMQPSLNP
1	1	l				GGEPV/ASDVVLLNHWKVTEF/
	Ì	l				SEVHRGDIVFIG\APKNPEQKIIK
i		l				RVICSLKGDIFQNHRTQKP\DVK
l	1	l			1	VPRGSHLGLEGDHSWDTVFDS
1	ı	1				NSFGPVSLGLLHAHATHILWPP
l	1	l				ERWQKLESVLPPERLPGTERRG
		l				MTA
23141	53509	Α	23275	90	664	NKYRLQPSPPRLALWDRASDPP
ŀ						APPEDPETAPAMEEDQELERKI
1	1					SGLKT\SMVEG\ERKTT\LELV\Q
		l		l		AAG\TDRP\CVTFVLHEEDHTLG
	1			l .		NFLRY\MIMKNP/EKVEFCGYT
	l			i		YG/HPSFQRAKLNLPHSRFRGYP
	ŀ					FPAVEPISREALNELM\NVC\QP
	ŀ					CAFTSFRASIKDYKESKSKQEM
	i	ŀ		1		ESTFLVLLWOFYKGGTVL
23142	53510	С	23276	108	320	` · · · · · · · · · · · · · · · · · · ·
23143	53511	В	23277	446	625	
23144	53512	A	23278	1	3042	
23145	53513	A	23279	1	3156	
23145	53513	A_	23279	[1	3156	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		i i		sequence		
23146	53514	Α	23280	518	2076	QQRR*MCQNLMTH\QNQRSGR
		1				SQNDF*RRRKNISERSAVYR*R
1		1			İ	QLYIRRQPYAERPS\HVEDKGH
		l				KYLVFEANTGTENGYQGEESLF
		ı				NKAYYGGGTNFFRKESQKLQQ
		l				SAKKRDAELANGALGIIELNND
1		!				YTLKKVMKPLITSNTVTDEIER
1		1			ł	ANVFKMNGKWYLFTDSRGSK
1		ı				MTIDGINSNDIYMLGYVSNSLT
		1				GPYKPLNKTGLVLQMGLDPND
1		1			l	VTFTYSHFAVPQAKGNNVVITS
		İ				YMTNRGFFEDKKATFAPSFLM
		l				NIKGNKTSVVKNSILEQGQLTW
		1				LOVAKRAGLGGGGSGRTVLRE
		l		1		RVRIEIASTHIALAVECVSVRV
1		l				WKVPRLPSRQKYAKHASQLVS
ł				-		NQKSEEEEYMKTFNISQQDLEL
1		1				VEVATEKITMLYEDNKHHVGA
		l				AIRTKTGEIISAVHIEAYIGRVTV
1		1				CAEAIAIGSAVSNGOKDFDTIV
į.		1	1			AVRHPYSDEVDRSIRVIPTYGT
İ		1				DEWEOWWNAFNEENLFCSEE
		l	1		1	MPSSDDEATADSQHSTPPKKKR
		1	}			KYIYVLVIAITMVMRFWQYING
23147	53515	A	23281	299	457	KTITYEVIJAINIVIJAA WQTING
23148	53516	A	23282	584	1185	QQRR*MCQNLMTH\QNQRSGR
231.0	55510					SONDF*RRRKNISERSAVYR*R
		1	1			QLYIRRQPYAERPS\HVEDKGH
		į .				KYLVFEANTGTENGYQGEESLF
1		l		1		NKAYYGGGTNFFRKESOKLOO
		l				SAKKRDAELANGALGIIELNND
1		1				YTSEKSNEAADHFKHVPOAKG
		1		l		NNVVITSYMTNRGFFEDKKATF
1		1				APSFLMNIKGNKTSVVKNSILE
		l				
23149	53517	-	23283		3618	QGQLTVN
23149	53517	A	23283	191	819	VALVECCED DICCTODUL FOCI *
23 130	33316	Ι^	23204	191	913	VNVTSFGSRPICSTSPVLFSGL*
1		1	1	l		GPVKEFDT*PSM*ISFELIPSIVIF
		l	1	I		EPRESVNKYHLPFILKTFARSISS
1		l	1	l		VTVFEVISGFITFFNV*SLFNSMI
				l		PRAPFANSASRFLALC*SFWLSL
1					ı	RKKFVPPP*YALLNKDSSPW/SS
			1		1	
						VFCSRVGFEYKVFMAFVFNVV

SEQ ID	SEO ID NO:	Mer	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
l			1	sequence		
23151	53519	Α	23285	2	2556	WRKIYQAN/GKTKKAGVAILVS
						DKTDFKPTKIKRDKEGHYIMVK
						GSIQQEELTILNIYAPNTGAPRFI
1			1			KQVLSDLQRDLDSYTIIMGDFN
						TPLSTLDRSMRQKVNKNTQEL
						NSALHQADLIDIYRTLHPKSTEC
1						TFFSAPHHMYSKIDHIVGCKAL
1						LSKFKRTEIITNCLSDHSAIKLEI
1						RIKKLTONCSTTWKLNNLLLND
1						YWVHNEMKAEIKMFFETNENK
						DTTYQNLWDTFKAVCRGKFIA
	ŀ					LNAHKRKQERSKIDTLTSQLKE
			ŀ			LEKQEQTHSKASRRQEITKVRA
1						ELKEIETQKTLQKINEFRSWFFE
			ĺ			RINTIDRPLARLIKKKREKNQID
		1	İ		İ	AIKNDKGDITTDPTETQTTIREY
		l		ł		YKHLYANKLENLEEMDKFLDT
	1	l	ŀ			YTLPRLNQEEVESLNRPITGSGI
		l			1	VAIINSLPTKKSPGPDGFTAEFY
	ľ			l		QRYKEELVPFLLKLFQSIEEEGI
				l	l	LPNSFYEASIILIPKLGRDIAKKE
			l		ŀ	NFRPISLMNIDAKILNKILAHRI
				l		QQHIKKLIHHDQVGFIPGMQG
	İ			l		WFDIRKSINVIQHINRTKDKNH
						MIISIDAEKAFDKIQQPFMLKTL
			l			NKLVLEVLVRAIRQEKEIKCIQL
	l					GKEEVKLSLFADDMIVYVENPI
						VSAQNLLKLISNFSKVSGYKIN
					ŀ	VQKSQAFLYTNNRQTESQIMSE
						LPFTIASKSIKYLGIQLTRHVKD
			ŀ	l	1	LFKDNYKPALNEIKEDTNKWK
1				1	i	NIPCSWIGRINIVKELEKTTLNFI
23152	53520	A	23286	I	3087	
23153	53521	Ā	23287	ı	741	MLGKKKPVCEVGGDSGVPKKE
		1		ľ		FOEGROVLFKLWRGOANKVG
				1		DNSIDSWKNAGRVFKDSDKFD
			İ		i	ANDPILKDQTQEWSGSATFTSD
				l		GKIRLFYTDYSGKHYGKQSLTT
	1			l	l	AOVKVSKS/V*HTQNORSGRSO
1	1	1	l	I		
	1			I		NDF*RRRKNIS*RSAVYR*RQL
				l		YIRRQPYAERPS\HVEDKGHKY
				l		LVFEANTGTENGYQGEESLFNK
1	l	1	l	I		AYYGGGTNFFRKESQKLQQSA
			[l		KKRDAELANGALGIIELNNDYT
						LKKVVKPLITSN

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			ŀ	sequence .		
23154	53522	Α	23288	849	1688	QQRR*MCQNLMTH\QNQPSGR
					1	SQNDF*RRRKNISERSAVYR*R
		l				QLYIRRQPYAERPS\HVEDKGH
		l				KYLVFEANTGTENGYQGEESLF
		l				NKAYYGGGTNFFRKESQKLQQ
i	1	l				SAKKRDAELANGALGIIELNND
į.		l				YTLKKVMKPLITSNTVTDEIER
		l				ANVFKMNGKWYLFTDSRGSK
		l				MTIDGINSNDIYMLGYVSNSLT
		1				GPYKPLNKTGLVLQMGLDPND
		1				VTFTYSHFAVPQAKGNNVVITS
		1				YMTNRGFFEDKKATFAPSFLM
		<u> </u>				NIKGNKTSVVKNSILEPGQLAV
23155	53523	A	23289	1	3171	MVKGSIQQEELTILNIYAPNTG
			1			APRFIKQVLSDLQRDLDSHTLI
	l	l				MGDFNTPLSTLDRSTRQKVNK
						DTQELNSALHQADLIDIYRTLH PKSTEYTFFSAPHHTYSKIDHIL
1		1				GSKALLSKCKRIEITNYLSDHS
		1				AIKLELRIKNLTQSRSTTWKLN
						NLLLNDYWVHNEMKTEIKMFF
		1				ETNENKDTTYQNLWDAFKAVC
		1				RGKFIALNAYKRKEERSKIDTL
		1				TSOLKELEKOEORHSKPSRROE
		l				ITKMRAELKEIETO
23156	53524	A	23290	1	3255	
23157	53525	A	23291	1	3924	
23158	53526	Α	23292	1	3139	
23159	53527	Α	23293	1	3210	MVKGSIQQEELTILNIYAPNTG
		l				ALRFIKQVLRDLQRDLDSHTIIM
1		l				GDFHTPLSTLDRSTRQKVNKDI
						QELNSALHQEDLIDIYRTLHPKS
	1	l				TEYTFFSAPHHTYSKIDHIVGSK
						ALLSKCKRTEIITNCLSDHSAIK
					i	LELRIKNLTQNRSTTWKLNNLL
						LNDYWVHNEMKAEIKMFFETN
						ENKDTTYQNLWDTFKAVCRGK
	1	1				FIALNAHKRKQERSKIDTLTSQL
						KELEKQEQTHSKASRRQEITKIR
	ļ	L				AELKEIETQ
23160	53528	В	23294	1	3127	
23161	53529	A	23295	1	3325	
23162	53530	Α	23296	11	3145	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuelcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amine acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nueleotide insertion)
23163	53531	A	23297	191	819	VNVTSFGSRPICSTSPVLFSGL*
						GPVKEFDT*PSM*ISFELIPSIVIF
		1		1		EPRESVNKYHLPFILKTFARSISS
						VTVFEVISGFITFFNV*SLFNSMI
				1		PRAPFANSASRFLALC*SFWLSL
				1		RKKFVPPP*YALLNKDSSPW/SS
		1				VFCSRVGFEYKVFMAFVFNVV
						RVSORMVVAGCIIAFIDKLLNV
						LICFSVSVKNRFVIFHSVDFECV
23164	53532	A	23298	I	3352	
23165	53533	A	23299	1841	2679	QQRR*MCQNLMTH\QNQRSGR
		l				SQNDF*RRRKNISERSAVYR*R
		l				QLYIRRQPYAERPS\HVEDKGH
		1				KYLVFEANTGTENGYQGEESLF
		l				NKAYYGGGTNFFRKESQKLQQ
		l				SAKKRDAELANGALGIIELNND
		l				YTLKKVMKPLITSNTVTDEIER
	1	1				ANVFKMNGKWYLFTDSRGSK
	İ	i				MTIDGINSNDIYMLGYVSNSLT
		1			1	GPYKPLNKTGLVLQMGLDPND
		1				VTFTYSHFAVPQAKGNNVVITS
				1		YMTNRGFFEDKKATFAPSFLM
						NIKGNKTSVVKNSILEQGQLTV
23166	53534	Α	23300	1207	1533	YVHCKPSGWTVRTFDKPRKRFI
						AFFIAGILFRAIKNHFLPRETLQ
						CLPYILTGFRRGQSEYFSIFSNM
	1	ı				DLADTVMFL*GAIRFSDLVNEQ
						IQHTFLIPGETICRLSEVV
23167	53535	Α	23301	I	3253	MGDFNTPLSTLDRSTRQKVNK
		l			1	DTQELNSALHQGDLIDIYRTLH
		l				PKSTEYTFFSAPHHTYSKIDHIV
		l				GSKALLSKWKRTEIITNYLSDH
		1				SAIKLELRIKNLTQSRSTTWKLN
		l	i			NLLLNDYWVHNEMKAEIKMFF
		l				ETNENKDTTYQNLWDAFKAVC
1		l				RGKFIALNAHKRKQERSKIDTL
	1	l				TSQLKELEKQEQTHSKASRRQE
l						ITKIRAELKEIETQKTLQKINESR
1	1	1				SWFFERINKIDRPLARLIKKKRE
			İ			KNQIDTIKNDK
23168	53536	A	23302	1	3095	
23169	53537	Α	23303	2	2678	

SEQ ID	SEQ ID NO:					Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
i	sequence		09/540,217	codon for peptide sequence	of peptide sequence	detection, (-possible nucleotide insertion)
				sequence		
23170	53538	Α	23304	1	3335	MVKGSIQQEELTILNIYAPNTG
						APRFIKQVLSDLQRDLDSHTLI
						MGDFNTPLSTLDRSTRQKVNK
1						DTQELNSALHQADLIDIYRTLH
						PKSTEYTFFSAPHHTYSKIDHIV
						GSKALLSKCKRTEIITNYLSDHS
						AIKLELRIKNLTQSRSTTWKLN
ľ						NLLLNDYWVHNEMKAEIKMFF
						ETNENKDTTYQNLWDAFKAVC
Ì						RGKFIALNVYKRKQERSKIDTL
						TSQLKELEKQEQTHSKASRRQE
						ITKIRAELKEIETQ
23171	53539	В	23305	1	2293	
23172	53540	Α	23306	I	3810	CLICCI I I I VIDED VICE II CD
23173	53541	Α	23307	255	1215	CVGQLAAAKR*RNSS*IQRLSR
						CVCSCGKPERR**HINLHVLSK
1						GRRQLN/QTAGKTRAVSLKTAI SSTPTIRS*KIRRKNGPVLOPLH
1						LTEKSVYSTLTIPTKAINTLYSK
						PTREOKTOTKAKNLYLTKRTT
		1			1	AAARTSSVKKARSFSRALKNA
1		1				MLS*RTAPSVS*S*IMITH*KK**
Ì		l				SR/CITSNTVTDEIERANVFKMN
						GKWYLFTDSRGSKMTIDGINSN
						DINMLGYVSNSLTGPYKPLNKT
		1				GLVLQMGLDPNDVTLTNSHFA
1		l	ĺ			VPQAKGNNVVITSYMTNRGFF
l		l				EDKKATFAPSFLMNIKGNKTSV
	1	l				VKNSILEQGQLTVN
23174	53542	A	23308	1	4539	
23175	53543	Α	23309	1	3229	
23176	53544	В	23310	1	3552	
23177	53545	A	2331I	1	3139	
23178	53546	Α	23312	I	3057	
23179	53547	Α	23313	I	3215	MGKKQNRKTGNSKTQSASPPP
		1				KERSSSPATEQSWMENDFDELR
1		1				EEGFRRSNYSELREDIQTKGKE
		1				VENFEKNLEECITRITNTEKCLK
		l				ELMELKTKARELREECRSLRSR
1				1	1	CDQLEERVSAMEDEMNEMKRE
						GKFREKRIKRNEQSLQEIWDYV
			1			KRPNLRLIGVPESDVENGTKLE
						NTLRDIIQENFPNLARQANVQIQ
1			1	1	1	EIQRTPQRYSSRRATPRHIIVRFT
		İ				KVEMKEKMLRAAREKDRSTRQ
	L					KVNKDTQELNSA

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23180	53548	Α	23314	I	1921	MVKGSIOOEELTILNTYAAHTG
						APRLIKQVLSDLQRDLDSHTIIM
				ŀ		GDFNTPLSTLDRSTRQKVNKDT
						QELKSALHQADLTDIYRTLHHK
						STEYTFFSAPHHIYSKIDHILGSK
						ALLSKCKRTEIITNYLSDHSAIK
				i		LELWIKNLTQNHSTTWELNNLI
						LNDYWVHNEMKAEIKMFFETN
						ENKDTTYHNLWDTFKAVCRGK
1		ĺ				FIPLNAHKRKQERSKIDTLTSQL
						KELEKQEQTN\HSKASRRQEITK
1						IRAELKEIETQKTL*KINESRSW
						FFERINKIDRLLARLIKKKREKN
						QIDAIKNDKGDITTDPTEIQTTIR
İ				l		EYCKHLYANKLENLEEMDKFL
				ĺ		DTYTLPRLNQEEVESLNRPITG
İ				l		AEIVAIINSLPTKKSPGPDGFTA
	1					KFYQRYKEELVPFLLKLFQSIEK
						EGILPNSFYEASIILIPKPGRDTT
i						KKENFRPISLMNIDAKILNKKL
1						AKRIQQHIKKLIHHDQVGFIPG
1			1			MQGWFNIRKSINVIQHINRAKD
						KNHMIISIDAEKAFDKIQQPFML
1						KTLNKL\GIKYLGIHLTRDVKDL
i						FKENYKPLLKEIKEDTNKWKNI
1		1		1		PCSWVGRINIVKMAILPKMCL*
		l				RRWTPLCLLTAAASLNRE*ERL
1						VQPLLWRQMYCGPRHCRQVPQ
		L_				HRRLNWSPSLRLS
23181	53549	B A	23315	584	2907 1278	OORR*MCONLMTH\ONORSGR
23182	33330	A	23310	384	12/8	SQNDF*RRRKNISERSAVYR*R
1		1				OLYIRROPYAERPS\HVEDKGH
i		l				KYLVFEANTGTENGYOGEESLF
1						NKAYYGGGTNFFRKESQKLQQ
						SAKKRDAELANGALGIIELNND
		1		1		YTLKKVMKPLITSNTVTDEIER
		1				ANVFKMNGKWYLFTDSRGSK
		1				MTIDVPOAKGNNVVITSYMTN
1		l				RGFFEDKKATFAPSFLMNIKGN
						KTSVVKNSILEQGQLTVN
23183	53551	A	23317	1	3345	K151 TKHSILLQOQLI YN
23183	10001	I.	23317	l <u>' </u>	3343	

			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23184	53552	Α	23318	2	3366	WRKIYQANGKQKKAGVAILVS
						DKTDFKPTKIKRDKEGHYIMVK
					l	GSIQOEELTILNIYAPNTGAPTFI
						OOVLSDLORDLDSHTLIIGDFN
						TPLSTSDRSTROKVNKDTOELN
						SALHOADLIDIYRTLHPKSTEYT
	-					FFSAPHHTYSKIDHILGSKALLS
	l					KCKRTEIITNYLSDHSAIKLELM
						IKNLTQNHSTIWKLNNLLLNDY
						WVHKEMKAEIKMFFETNENKD
	İ					TT\YQNLWDTFKAVCRGKFIAL
						NAHKRK\QER
23185	53553	Α	23319	1	2611	MGDFNTPLSTLDRSTRQKANK
						DTQELNSALHQVDLIDIYRTLH
						PKSTEYTFFSAPHHTYSTTDHIL
						GSKALLSKCRRTEIITNYLSDHS
	İ					AIKLELRIKNLTQNRSTTWKLN
					1	NLLLNDYGVHNEMKAEIKMFF
	l					ETNENKDTTYQNLWDTFKAVC
	ĺ					RGKFIALNAHKRKQERSKIDTL
		ļ				TSQLKELEKQEQTHSKASRRQE
	İ					ITKIRAELKEIETQKTLQK\INES
						RSWFFERINKIDRPLARLIKKKR
		1				EKNQIDAIKNDKGDITTDPTEIQ
						TTIREYYKHLYANKLENLEEM
						DKFLHTYTLPRLNQEEVESLNR
						PITGSEIVAIINSLPTKKSPGPDG
						FTAEFDQRYKEELVPFLLKLFQ
						SIEKEGILPNSFYEASIILIRKPGR
						DTTKKENFRPISLMNTDAKILN
						KILAKRIQQHIKKLIHHDQVGFI
						PGMQGWFNIHKSINVIQHINRA
						KDKNHMIISIDAEKAFDKIQQPF
						MLKTLSKLGIDGTYFKIIRAIYD
		l				KPTANIILNGQKLEAFPLKTGTR
		l				QGCPLSPLLFNIVLEVLARAIRQ
				ļ		EKEIKLISNFSKVSGYKINVQKS
	1		l			QAFLYTNNRQTESQIMSQLPFTI
						ASKRIKYLGIQLTRDVKDLFKE
	1		l	l		NYKPLLKEIKEDTNKWKNIPCS
	l		l			WVGRINIMKMAILPKVIYRFNA
	1		l			IPIKLPMTFFTELEKTTFKFIWN
		l		l		QKRARITKSILSQKNKAGGITLP
	l	ı	ı	l	ı	DFKLYYKATVTKTA WYWYQN

SEQ ID	ISEO ID NO:	Mot	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	1	
		L				
23186	53554	Α	23320	1	3099	MGELITPLSTLDRSTRQKVNKD
						TQELNSALHQGDLIDIYRTLHP
				l		KSTEYTFFSAPHHTYSKIDHILG
						SKALLSKCKRTEIITNYLSDHSA
					i	IKLELRIKNLTQNRSTTWKLNN
1				l		LLLNDYWIHNEMKAEIKMFFET
				l		NENKDTTYQNLWDAFKAVCR
ŀ		İ		l		GKFIALNAHKRKQERSKIDTLT
					1	SQLKELEKQEQTHSKASRRQEI
				l		TKIRAELKEIETQKTLQKINESR
						SWFFERINKIDRPLARLIKKKRE
					ľ	KNOIDTIKNDK
23187	53555	A	23321	911	1539	VNVTSFGSRPICSTSPVLFSGL*
					ŀ	GPVKEFDT*PSM*ISFELIPSIVIF
l					1	EPRESVNKYHLPFILKTFARSISS
					Ì	VTVFEVISGFITFFNV*SLFNSMI
Ī						PRAPFANSASRFLALC*SFWLSL
				l		RKKFVPPP*YALLNKDSSPW/SS
						VFCSRVGFEYKVFMAFVFNVV
						RVSORMVVAGCIJAFIDKLLNV
						LICFSVSVKNRFVIFHSVDFECV
23188	53556	Α	23322	191	820	VNVTSFGSRPICSTSPVLFSGL*
						GPVKEFDT*PSM*ISFELIPSIVIF
						EPRESVNKYHLPFILKTFARSISS
ŀ						VTVFEVISGFITFFNV*SLFNSMI
						PRAPFANSASRFLALC*SFWLSL
				1		RKKFVPPP*YALLNKDSSPW/SS
				1		VFCSRVGFEYKVFMAFVFNVV
						RVSORMVVAGCIJAFIDKLLNV
1	į.	l			i	LICFSVSVKNRFVIFHSVDFECV
23189	53557	A	23323	965	5050	TWKGTTSTSRCKIMPKYRSTRO
23,05	3333,	ľ	23323	,00	5050	KVNKDTQELNSALHQADLIDIY
		l				RTLHTKSTEY/TFFSAPHHTYSK
1		l				IDHIVGSKALLSKCKRTEIITNY
1		l		l	ŀ	LSDHSAIKLELRIKNLNQSRSTT
	i			ŀ		WKLNNLLLNDYWVHNEMKAE
ŀ						IKMFFETK\ENK\DTTYQNLWD
1						AFKAVCRGKFIALNAHKRKQE
1	l		l	1		RSKIDTLTSQLKELEKQEQTHS
1	l		l	l		
1						KASRRQEITKIRAELKEIETQKT LQKINESRSWFFERINKIDRPLA
	l	ı		l		RLIKKKREENO
23190	53558	A	23324	1	2787	RLIKAKEENŲ
23190	53559	A	23324	1	902	
	53560	A	23325	1	3640	
23192	133200	IA.	23320	1	3040	

SEQ ID	SEQ ID NO:					Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23193	53561	Α	23327	1	3686	MHIVVETALSASWQNKAKPPA
						RVLLQVVPNVWFLVAVVWEL
						YPSLDLMDRSIECSSSPATEQS
		[WTENDYDKLREEGFRRSNYSE
						LQEDIQTKGKDVENFEKNLEEC
						ITRITNTQKCLKELMELKTKAR
		1				ELREECRSLRSRCDQLEERVSV
		1				MEDEMNEMKQEGKFREKRIKR
						NEQSLQEIWDYVKRPNLRLIGV
						PESDGENGTKLKNTLQDIIQENF
		1				PNLARQAKVQIQEIQRMPQRYS
						LTRATPRHIIVRFTKVE
23194	53562	Α	23328	1	3433	MGDFNTPLSTLDRSTRQKVNK
						DTQELNSALHQADLIDIYRTLH
1						PKSTEYTFFSAPHHTYSKIDHIL
						GSKALLSKCKRTEIITNYLSDHS
						AIKLELRIKNLTQNRSTTWKLN
						NLLLNDYWVHNEMKAEIKMFF
						ETNENKDTTYQNLWDAFKAVC
						RGKFIALNAHKRKQERSKIDTL
						TSQLKELEKQEQTHSKASRRQE
İ						ITKIRAELKEIETQKTLQKINESR
1						SWFFERINKIDRPLARLIKKKRE
						KNQIDAIKNDK
23195	53563	Α	23329	1	3070	MEGEMNEMKREGKFREKRIKR
						NEQSLQEIWDYVKRPNLRLIGV
						PESDGENGTKLENTLQDIIQENF
		1				PNLARQANVQIQEIQRTPQRYS
						SRRATPRHIIVRFTKVEMKEKM
1						LRAAREKDRSTRQKVNKDTQE
						LNSALHQADLIDIYRTLHLKSTE
						YTFFSAPHHTYSKIDHILGSKAL
						LSKCKRTEIITNYLSDHSAIKLE
						LRIKNLTQNRSTTWKLNNLLLN
		6				DYWVHNKMKAEIKMFFETNEN
		L_				\KDTTYQNLWDA

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nuelcotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	eodon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	eodon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequenee		
23196	53564	I .	23330	3	2372	TTNCLSDHSAIKLELRIKKLTEN
23196	33304	A	23330]3	2312	
		l		l		HTTTWKLNNLLLNDYWVDNE
		ĺ				MKAEIKMFFEINEKK\DTTSQNI
l				l		WDTFKAVCRGKLIALNAHKRK
				l		QERSKIDTLTSQLKK\LEKQEQR
						NSKASRKQEITKIRAELKEIETQ
1					l	KKTLQKINESRSWFFEKINKIDO
ŀ						PLARLIKKKREKNQVDAIKNDK
1					l	GDIITDPTEMQTTIREYYKYLY
l				l		ANKLENLEEMDKFLNTYTLQR
		ŀ				LNQEGIESLNRPITGSEIEAIINSL
İ						PTKKSPGPDGFTAKFYQRYKEE
l		l				LVYPEPRSESECLSNIREFLRGC
		[GASLRLESIREDRNGRSQKTVH
l						TEGDMNMNIKKIVKQATVLTF
						TTALLAGGATQAFAKENNOKA
						YKETYGVSHITRHDMLQIPKQQ
i		1				QNEKYQVPQFDQSTIKNIESAK
1		1				GLDVWDSWPLONADGTVAEY
1					Į.	NGYHVVFALAGSPKDADDTSI
1						YMFYQKVGDNSIDSWKNAGR
		1				VFKDSDKFDANDPILKDQTQE
ļ.		1				WSGSATFTSDGKIRLFYTDYSG
		1		ł		KHYGKOSLTTAOVNVSKSDDT
l		1				LKINGVEDHKTIFDGDGKTYON
		1				VOOFIDEGNYTSGDNHTLRDPH
		1				YVEDKGHKYLVFEANTGTENG
1						YQGEESLFNKAYYGGGTNFFR
		1		1		KESQKLQQSAKKRDAELANGA
1		1				LGIIELNNDYTLKKVMKPLITSN
1		1		1		TVTDEIERANVFKMNGKWYLF
l				1		
		_				TDSRGSKMTIDGINSNDIYMLG
23197	53565	A	23331	2	2987	APRFIKQVLSDLQRDLDSHTIIM
l		1		1		GDFNTPLSTLDRSTRQKVHKDT
		1		i		QELNSALHQADLIDIYRTLHPK
		1				STEYTCFSAPHHTYSKIDHIVGS
1		1			1	KALLSKCKRSEIITNCLSDHSAI
		1			i	KLELRIKKLTQNRSTTWKLNNL
		1				LLNDYWVHNEMKAEIRMFFET
		1				NENKDTTYQNLWDTFKAVCRG
						KFIALNAHKRKQERSKIDTLTS
		1				QLKELEKQEQTHSKASRRQEIT
		1				KIRAELKEIETQKTL\QKINE\SR
		l				SWFF\ERINK
23198	53566	Α	23332	339	701	PDKEGNEIWVDMYTVKPSGWT
[1	1		1		VRTFDKPRKRFIAFFIAGILFRAI
		1	l	1		KNHFLPRETLOCLPYILTGFRRG
	1	l				QSEYFSIFSNMDLADTVMFL*G
		1		I		AIRFSDLVNEQIOHTFLIPGETIC
		1				RLSEVV
23199	53567	A	23333	190	502	The state of the s
23199	22307	ΙΛ.	23333	170	302	

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	ł	
23200	53568	Α	23334	4472	8032	KKSGNNRLMRVHGKKRFLWQ
						KQGVCEVEIQAQRESARDGTD
		1		ŀ		REDAESPHAEVTVPRGTRTEILT
				l		LWASRDRPASACGVTVPKGST
				l		THPGGMDRVCGVDTAVLHWM
				l		KQVQGISNLEVETGPRRRDMD
						EAGSRHSQQTNIGTENQTPHVL
				l		THKWQLNRNSACDKLHKPHSV
		l				PGGVAAGDVGGKTISNPGVCAI
						EYQVDGSPHIWEDVNKGVYIEE
				l		DRAPLMERSSSPAMEQSWMEN
						DFDELREEGFRRSNYSELRED
23201	53569	Α	23335	1	2374	MVSISDLVICPPRHPKVLGLQGP
						PGLDSISDPSAGAGFLDWGEIG
	ŀ			l		MPGPGRAGHQALCKCDCQCLE
		l				KTTTKAPGKMPKSTRSGPVRV
		l		l		RLADGPNRCAGRLECGMPDAG
				l		EQCVMTTGTSGRHCGLLGTGL
		1				WKGYTDLTIIPPGPGTPPQERTC
		1				QGDYHSGGTWTHSPLETTRRP
		l			{	GSSSPAIRRLPAQMLLLPARPPH
	i					PRSSSPEAMDPPPPKAPPFPKAE
			l		1	GPSSTPSSAAGPRPPRLGRHLLI
		l				DAN/GVYPYTYTVQLEEEPRGP
						PQREAPPGEPGPRKGYSCPECA
					l	RVFASPLRLQSHRVSHSDLKPF
						TCGACGKAFKRSSHLSRHRATH
						RARAGPPHTCPLCPRRFQDAAE
						LAQHSWGTPRGPLLAAACNCE
						VARGRLESPGPERLLHGYGGRE
						EEGGWGRAAGGLDRVEGFISS
						KAHHYLLIDTQGVPYTVLVTRS
		1		l		HRGSQGPVGLQARKVLQLPRV
					1	LKGLRVHVHLQRHSITHSEVPQ
						DFAGSLDSFQTPGESLRLVFRA
						LDTTQSSRISKAEPCLKEEPLSL
				i		GDLPYMHTTLCFCRKRRASPGP
					1	GTLQRGALAWPDWASPRALPV
	1			I		PSLSSTTRSPAAPLFAVPLSGRT
	l		1			TQAMAFDGIIFQGQSQRSAGLT
	1			I		TTSRFLACQRPLRLCAWWASRS
				I	1	PRCTLRRPVGLRPGVHPRPRLV
		1		l		YRDLKPENVMASGOPRDRPOP
	I	l	1	I		WFAWPPRPTRFCGGCWTLTPK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codnn, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23202	53570	A	23336	853	1602	PGAAWSRPDLRGCCTEPQPALR
23202	33370	<u> </u> ^	25550	055	1002	MLVLASPCPOPLAFSSVETMEG
i		ı				PPRRTCRSPEPGPSSSIGSPOASS
						PPRPNHYLLIDTQGVP\YTVAW
1						LDEESOREPGASGAPGOKKCYS
						CPVCSRVFEYMSYLQRH\RFSH
1						SEVKPFE\CDICGK\AFKRSQHL/
1						SARHHSIHLGGWVGGPTACRFC
						PRRF\RDAGELAQHSRVHSGER
	İ					PVSVSTLPSPLYGAEHTAETHA
						VEASHEAGLPGAPGTHRTLQG
		1				ALDSCPDTW
23203	53571	A	23337	1	1144	MGHHLMIDKLVALGGLYYAIQ
						RHYATKCSVLKNDQILVIGLFM
1						IQNVIYRKHFANPLSALFLQQGI
						ELFAAIAEIHITTVAERNHAITQI
						RLEAQTIFDSLKEWHNAIRKSL
						DYEALQAFEFRVGATDRGFPAL
						SSEALVRVLVLDANDNSPFVLY
		l				PLQNGSAPCTELVPRAAEPGYL
		1				VTKVVAVDGDSGQNAWLSYQ
		ŀ				LLKATEPGLFGVWAHNGEVRT
					l	ARLLSERDVAKHRLVVLVKDN
						GEPPRSATATLQVLLVDGFSQP
						YLPLPRAAPAQAQADSLTVYL
						VVALASVSSLFLFSVLLFVAVR\
						LCRRSRAAPVGRCSVPECPFPG
	l	l				HLVDVSGTGTLSQSYQYEVCLT
						GGSGANEFKFLKPVIPNLLSRDS
23204	53572	Α	23338	1	1260	MVCYIQENLPFLLKPSVENFYIL
						ITEGALDREIRAEYNITITVTDL
						GTPRLKTEHNITVLVSDVNDNA
		1				PAFTQTSYTLFVRENNSPALHIG
					1	SVSATDRDSGTNAQVTYSLLPP
1						QDPHLPLASLVSINADNGHLFA
						LRSLDYEALQAFEFRVGATDRG
1		1				SPALSREALVRVLVLDANDNL\
						PFVLYPLQNGSAPCTELVPRAA
		1				EPGYLVTKVVA VDGDSGQNA
		1				WLSYQLLKATEPGLFGVWAHN
1			1	1	l	GEVRTARLLSERDAAKHRLVV
	1	1	1		l	LVKDNGEPPRSATATLHVLLVD
1						GFSQPYLPLPEAAPAQAQAEAD
			1	1		LLTVYLVVALASVSSLFLLSVL
		1		1		LFVAVRLC\RRSRAASVGRCSM
			1	1		PEGPFPGRLVD\VSGTGTLFQSY
						QYEVCLTGGSETGEFKFLKPITP
	L		L	l		HLPPHRGGKEIEENSTLPNSFGF

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X-Unknown, *=Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
23205	53573	A	23339	123	5794	SSSDSGSLKGACTEVKMEPAGE RFPEQRQVLILLLLLEVTLAGW EPRRYSVMEETERGSFVANILAN DLGLGVGELAERGARVVSEDN EQGLQLDLQTGQLLINEKLDRE KLGGFTEPCIMIFQVLLKKPLE VFRAELLVTDINDHSPEFPERE MTLKIPETSSLGTVFPLKKARD LDVGSNNVQNYNISPNSHFHVS TRTRGDGRKYPELVLDTELDRE EQAELRILITAVDGGSPRSGT VOILLIVLANDNA
23206	53574	C	23340	20	313	
23207	53575	Α	23341	119	247	RSRCISASPRRPASFRKSIPEEED ESLNI*KGQVSILFFYIK
23208	53576	В	23342	55	288	
23209	53577	A	23343	1	3576	MAFLFEFSDLVSLLSIGTMLAY SLVAFSVLVLRYQPDENFSKNE KPKEEVVBMNSVPKAESPACVP EASSTPASLWSPVSTIPTPRWGR IVYGCAFILVVLLSMLCLVLAH WPKRLFSGELIVIAAAVLLLVLI VGFTFTVWRQPQSNTFLVFKEP HISTSQNLSLPEHPMEYLLRTGI EPDVRGVLTVNALPAGPPVACS PLVSISAHVYLMMQMTTETWA QCGVWMLIALECPNLKLKKPP WLIMPLAMTMYA

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SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X-Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
23210	53578	A	23344	3	1762	SVREOFGHIVRVRFFSI.SPLPLR PWPULCPAR.RTMGSILSRRIAG VEDIDIQANSAYRYPPKSGNYF ASHFFMGGEKFDTPHEGHLIFG ENMDLNFILGSRPSQFPYVTPAP HEFVETTLRSI,VNIRKDSLRLVR YKDDADSPTEDGDKPRVLYSL EFTFDADARVAITIYCQASEEFL NGRAVYSFKSPSLQSETVHYKR GYSQQVSLPSFKIDFSSWKDDE LNFDLDRGVFPVVIQAVVDEG/ DSVVEVTGHAHVLLAGFEKHM DGSFSVYRLEXDGKOVDRVSYLL QEIYGIENKNNQETKPSDDENS DNSHECVVCLSDLRDTLLFC HLCLCTSCADTLRYQANNCPIC RLPFRALLQIRAVRKFOALSP YSSPVLAGSLEHDEHSGPFKK SKPHFASLASKKPKRETNSDSV PPGYEPISLLEALNGLRAVSPAI PSAPLYEETIYSGISDGLSQASC PLAAIDHILDSSRQKGRPQSKAP DSTLRSYSSPHEEDEEKI.SEDV DAPPPLGGAELALRESSPESFI TEEVDESSSPQGTRAASIENVL QUSSPEHLGGRGPPAANSIENVL QUSSPEHLGGRGPPAANSIENVL QUSSPEHLGGRGPPADIVLPALG
23211	53579	A	23345	35	3134	RPREGPTWPQLPGSSPRRLEPW SPGSSSRGLFPWSPGSSSRGLF WSPGSSSRGLFPWSPGSSSRGLF PWSPGSSSRGLFPWSPGSSSRG GSSRGHPGPPPGGSSRG GSSRGHPGPPPGGSSRGHPGPP GALPVVTRVLLPGLFPWSPGSSSRGHPGPPWSPGSSSRGLFPWSPGSSRGHPGPPWSPGSSRGLFPWSPGSSRGHPWSPGSSRGHPWSPGSSRGHPWSPGSSRGHPWSPGSSRGHPWSPGSSRGHPWSPGSSRGHPWSPGALPVVTRVLLBGTPWSPGSSRGHPWSPGALPVVTRVLLBGTLPVVTRVLLBGTLPVVTRVLL
23212	53580	A	23346	3	91	7.11.122
23213	53581	A	23347	3	174	
23214	53582	A	23348	3	360	
23215	53583	В	23349	158	368	

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	eodon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		***
23216	53584	A	23350	3786	4928	PPPARTRPCASLTPEKAKWWRS
		1				KPGLTRAPARCGLSSPOTGSCS
	1	1		1		APASAG*VSGNSRSGTRRGLRP
		l				TRG*GPCGPSSRARAISSPRAS/L
		l				RMSQRELGLWDPNNFEEPVAL
						QEMDTSNGVLLPFYDPD\$SIVY
		1				LCGKVLTAGQGEQGTGWRCG
		1				GPCPGAPLNRLILQGDSSIRYFE
		1				TDEPPFVHYLNTFSSKEPORGM
		1				GFMPKRGLDVSKCEIARFYKLH
		l		i	Ì	ERKCEPHMTVPRKSDLFQDDL
						YPDTPGPEPALEADEWLSGQD
					İ	AEPVLISLRDGYVPPKHRELRV
						TKRNILDVRPPSGPRRSQSASDA
						PLSVRSALLHSGPIYISPNRPYTT
				l		RPLCPPPQQQHTLETLLEEIKAL
						RERVQAQEQRITALENMLCELV
23217	53585	Α	23351	1	1898	MSRKGPRAEVCADCSAPVGPD
						FALAISELVSPDIFMQSHSENAIS
				ì		VKEIVTEIESISQGVGQIQLKGDI
i						LPNPCHTPKKNSIHELLLERAQT
l		i				PENKPGHMEQDEDSCTAQPEL
						AKDSGMCNPEGCLTTHSSIADL
				ľ		EEGEQLRGNKSSRAQGCTQNS
İ				į.		KNDSSVADLAPKGKSDEAPPEH
		1		1	1	SFVLKEPEMSKGKGKYSGSEA
1		1				GSLSHSEQNATVPAPRVLEFDH
						LPDPQEGPGPEASNGLKVSQKS
1						EGSQPSSSRSPLEFLKEAGVPNQ
ł		1				TLSSGTPDPYFMALLGPKLNSR
						RPHLHPVSHTLDLGNSPPLPQQ
						FAKCQRGRKGPEGKYPLHYLV
				i		WHNRHRELEKEVRAGQTLITQ
		1				TLSQANPTAITAEEYFNPNFELC
						NRDMGRPMELTTKTQKFKAKL
		1		i		WLCEEHPLSLCEQVAPIIDLMA
l		1		1		VSNALFAKLRDFITLQIPIFHILN
1		1		1	1	ARITFGNLNGCDEPVPSVRGSPS
ļ.		1		1	}	SETPSPGSDSSSVSSSSSTSEAPR
	1		1			ENACPSALPGVASCRGCEISPAL
						FEAPRGYSMMGGQREAATRDD
1						DDDLLQFAIQQSLLEAGSEYDQ
1	1	1	1		[VTIWEALTNSKPGTHPMSYE/R
1					1	SPTGQGQVPAATCSG/RYDEQL
1		1				RLAMELSAQEQEERRRRARQE
		١	00000			EEELERILRLSLTEQ
23218	53586	Α	23352	341	711	

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ı	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		ł		sequence		
		ᆫ				L
23219	53587	Α	23353	57	719	TSDLMAEIIQERIEDRLPELEQL
		1				ERIGLFSHAEIKAIIKKASDLEY
		1	1			KIQRRTLFKEDFINYVQRIGYSF
		1				KKDEIENSIVHRVQGVFQRASA
		1				KWKDDVQLWLSYVAFCKKWA
		1				TKTRLSKVFSAMLAIHSNKPAL
i		l				WIMAAKWEMEDRLSSESARQL
i		1				FL\RALRFHPECPKLYKEYFRM
		1				ELMHAGKTGGRRRKNLKKPV
		l				WMWRILIILKKSLRASWHGSST
23220	53588	A	23354	27	352	TESVVTIMSLLNKPKSEMTPEE\
23220	33300	ľ	25554	["	1552	REEEFNTGPLSVLTQSVKNNT
		l				OVLINCRNNKKLLGRVKAFDR
		l				HCNMVLE\NVKEMLTEG/DRYI
		l				
23221	53589	A	23355	32	575	SKMFL\AGDPVIVVLRNPLIAGK
23221	133389	A	23333	32	373	STRLDLPKCWDYRSLFNKP\KS
		1				EMTP\EELQKREE\EEFNT\GPTP
	l	l				LCLTQS\VKNNTQSAQSTC\RNN
		l				KKLPGAA*RPFDRASPPCSPPGT
Į.	ŀ	l				CNMVL\ENVK\EMWT*GTQEV
	ŀ	l				GKGKEGSSKP\VNKDRLTSSKM
		l				FPGGGGFSSFVVPCGNPFHSAG
		l				QIRGPPVLFDKNSHPLVLLKTA
						CHLVFENY
23222	53590	Α	23356	103	713	VTSGCGKKKVWLDPNET\NEIA
i	l	l		1		NANFRSADPGRLIKDGLIIRKPV
		1		1		T\VHSRARLPEKTPLARCKGHG
						TLGIGKRKGTANA\RMPKEGHH
		l				GMRRMRILAPACSRRYRESKKI
						D\RHMYHSLYLKVKG\NVFKNK
	l				1	ADFSWEHIHKAEGRARPRKEAS
		1	ŀ		1	WLDQAEA\RRSKDQGKHGKRR
	i					EERLPGORKEEINORLYSKEGR
						RPRNKNLTL
23223	53591	Α	23357	1	2847	
23224	53592	Α	23358	I	2823	
23225	53593	Α	23359	3	2858	AMFGFQRRGLGTPRLQLWLLL
		1				LEFWEVGSGOLHYSVSEEAKH
1	l	l	1		1	GTFVGRIAQDLGLELAELVQRL
1	1	1				FRVASKTHGDLLEVNLONGILF
1		l				VNSRIDREELCGQSAECSIHLEV
1		l	1			IVDRPLOVFHVNVEVKDINDNP
1		l				PVFSLREQKLLIAESKQSDSRFP
1	l	l		l		LEGASDADIEENALLTYRLSKN
1	1	1		I	i	
						EYFSLDSPTNGKQIKRLSLILKK
	1	l	1		l	SLDREKTPELNLLLTATDGGKP
1		I		1		ELTGTVRLLVQVLDVNDNDPE
L	L	L_		l		FDKSEYKVSLMEN

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23226	53594	A	23360	194	931	NSPVVPVMASGQFVNKLQEEVI CPICLDILOKPVTIDCGHNFCPO
						CITQIGETSCGFFKCPLCKTSVR\
					ŀ	RDAIRFNSAVAEIWWEKIQALQ
					İ	ASEVQSKRK\EAT\CPRHQEMF
						HYFCEDDGKFLCFVCRESKDH
						KSHNVSLIEEAAQNYQGQIQEQ
						IQVLQQKEKETVQVKAQGVHR
						VDVFTDQVEHEKQRILTEFELL
						HQVLEEEKNFLLSRIYWLGHEG
						TEAGKHYVEIPLMPTVERSQEA
23227	53595	Α	23361	2	381	AINWTSARPVSQCWDP\CPPSQ
						VVCLAGLPP*RVCQNPVKGP\N
						CPLAHTSEPTGHCLL*TRGMCF
						PSLCGQLCP*SGPPDCPSPRALH
						HTGFCPVALLVAESQVRARAC
						GPEPHRAGRASRWCLHAG
23228	53596	A	23362	213	574	PQTHTGNHSERGRQPITRSPEKP
						EP\CPSKRAPQTQQP\PWAPSLC
						PPALGPCKGTRRPRVQVHRPTS
						SQKCPLPAWFHHLPLWPTSAAR
						SSSTETPTGQQMRKLRHPAAPR
						SANALSLGE_
23229	53597	Α	23363	21	456	
23230	53598	Α	23364	2	98	
23231	53599	Α	23365	64	241	PWPLASNPRGCLP*PSTLN*PTP
		l	l	l		SSQPGMSLDSLDRHPP*TRETAS
						GIGGQRPGSDPARTPHFISQTQ
23232	53600	Α	23366	209	353	

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SEQ ID	Toro In NO.	124.4	LOCO ID NO.	Ter to all do	In	I
NO:	of peptide	Met	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence	1.00	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23233	53601	Α	23367	1766	3292	SSCPTTRVLIMKGAGHPCYLDK
		l				PEEWHTGLLDFLQGLQLWLMH
		l	l			MOQLMSLAVVSGRLSRGQCGT
		l				ERPSRRAAAAAAAGOPORGHF
	1					OHIEERLETNVONCOSPDTLCP
	İ	l				THCSPVLNEPCPEVSVVAVAPA
		ĺ				SPDRMSGSDGGLEEEPELSITLT
		ļ				LRMLMHGKEVGSIIGKKGETV
		1				KRIREOSSARITISEGSCPERITTI
		1				TGSTAAVFHAVSMIAFKLDEDL
		1				CAAPANGGNVSRPPVTLRLVIP
		1				ASOCGSLIGKAGTKIKEIRETTG
		1				AQVQVAGDLLPNSTERAVTVS
		1				GVPDAIILCVRQICAVILEGFSV
		1				QGQYGAVTPAEVTKLQQLSSH
		1	1	ı		AVPFATPSVVPGLDPGTOTSSO
		1	}			EFLVPNDVSRGWDGWRLIGCVI
		1				GRQG\TKINEIRQMSGAHIKIGN
		ı				O\AEGAGERHV\TITGSPVSIALA
		l				QYLITACLETAKSTSGGTPSSAP
		l				ADLPAPFSPPLTALPTASPLAC
		l				WAHPMPSPSPTLIGFKAHALLG
						FTTCFPRAAAGLGGLHRODGSS
23234	53602	В	23368	1	363	
23235	53603	A	23369	486	947	LDGAGLLAPVKAHLOKKGPRL
		1				GTRGCAMPRLGSTPFLKGQCL
		l			1	ANSRNTLEAVSLTESSASRVPP
		l				APVLGSRRPHSPSPWP/P/YYTP
						VPPPPRMALVPPNMHGQGTVG
						LTSVPLACTVAMLGSYKQAGA
						FGPLANCYGGLGHAQCWGQEF
		1				СРТОСР
23236	53604	A	23370	38	680	GSRLRRLOAAAARPALPLPLPP
						WEWKHLPHVPEAKWWLTTAR
1		t				HSAAYRADPLRVSSRDKLTEM
1		İ				AASSOGNFEGNFESLDLAEFAK
ĺ			ŀ			KOPWWRKLFGOESGPSAEKYS
						VATOLFIGGVTGWCTGFIFOKV
						GKLAATAVGGGFFLLQLANHT
						GYIKVDWQRVEKDMKKAKEQ
		1	1			LKIRKSNQIPTEVRSKAEEVVSF
						VKKNVLVT\GGFFGGFLLGMAS
23237	53605	С	23371	30	200	
23238	53606	Ā	23372	14	161	
23230	122000	<u>'``</u>	20012		101	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	ĺ			sequence		
23239	53607	Α	23373	69	715	APALPGCEHMMAIRELKVCLL
		l		1		GDTGVGKSSIVWRFVQDHFDH
		1				NISPTIG\ASFMTKTVP\WENDFT
		l				KFLIWDTAGQERFSFIGLPLYYS
		l				RLQLAAVYSCYGYLPSRDFIFIT
		l				LKE\WV\KRLKELGPE\NIVMAI
		1				AGNKCDLSDI\REVPPEGMLKE
		1				Y\AESIGAIVVETSAKNAINI\EE
ŀ		1	į			LFQGIS\RQIPPLDPPMKNGKQW
						DHOKLRSOPMOASRRCC
23240	53608	A	23374	3	293	VTSTMAYRAQGQKV\QKVMV
						QPINLIFRYLQNRSRIQVWLYEQ
l		1				VNMRIEGCIIGFDEYMNLVLDD
		1		1		AEEIHSKTKSRKOLGRIMLKGD
		l				NITLLQSVSN
23241	53609	A	23375	3	393	HESCSQRQRAGCALCENSHPW
		ļ.,		ľ		RSRGPGVQKSAEGLWCOPICRT
		l	İ			SSSRYL\QNRIAGFQVWLYEQV
l		l				NM\RI\EGCIIGF\DEYVNLVLDD
		ì				AEEIHSKTKSRK\OLGR\IMLK\G
		1				DNITLLQTCLPPRMVHGSGEIV
23242	53610	С	23376	129	425	
23243	53611	Α	23377	470	635	ENSSSHHIIKPFSSGCSIHGFSGH
		1				INVTSVTIL*PIKSCHIRCHF*CA
						QYMVADN
23244	53612	Α	23378	7I	566	PLFPPCLSPVVAGWARGHMLG
				1		QSHQQVAAPLKHRPITPTELCQ
		1				KCLSLIKKTKFLNAYITVSEEVA
						LKHAEESEKRYKNGQSLGDLD
l		1				GIPIAEKDNFSTSGIETT\CASNM
		ı				LKGYIPPYNATVDQKLLDQGA
ĺ						LLMGKTKLNDFAMGSGSTDGV
						FGPVKNPWSYSK
23245	53613	Α	23379	255	1042	ILAARMGKQNSKLRPEVMQDL
1	l	1	ļ	l		LESTDFTEHEIQEWYKGFLRDC
	l	1		l	1	PSGHLSMEEFKKIYGNFFPYG\D
1		1	l	1		ASKFAEH\VF\RTFDANGDG\TID
	1	1	l	l	l	FREFIIALSVTSRGKLEQKLKW
	l	1			i .	AFSMYDLDGNGYISKAEMLEIV
ŀ	ŀ	1				OAIYKMVSSVMKMPEDESTPE
1	l .	1	l		l	KRTEKIFROMDTNRDGTWVPH
ŀ		1	l			LALIYRTAEGGFAVORASVSFY
1		1	l	l		PPPSAGNAMVNEAAVFLEIPLG
1		1	l			SSPGOVSAVCHSGRKLSLEEFIR
		1	l			GAKSDPSIVRLLQCDPSSAGQF
	l			L	L	G. H.C.D. G. T. C.D.QCDI GG/TOQ1

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nuclcotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	1			sequence		
23246	53614	A	23380	1	721	MDGNLRKIRFEKHAITEEFRTT
23240	33014	^	23300	l'	1/21	EPGLAALRLPSTWIRGVRTSES
	İ	l				KREVGKGGOSOOYENRDTEGE
	l	l				GHVMTKARDWSKAAASQRLP
	1	ŀ			i	
						ANHLKLRLCIDGFRSHSLPEEGP
1		l				KKPCQHHAAQEEPCTRSGRLPQ
		l				AHEGSISDCCTSSEQGSMDIGSV
1		1			İ	EPGTGYNLLVCHLL/STLEKRSI
		1				WAGVSRFSRYHLSRLPLARK\G
		1				NPLTPCASQLLLPLPSTTQKRQS
						GHALYSFPYRPQPKNDNLVMP
23247	53615	Α	23381	80	562	DRAMWGDLWLLPPASANPGT
	1					GTEAEFEKAAEEVRHLKT/RSPS
		1				DEEMLFIYGHYKQATVGDINTE
	i .					RPGMLDF\TGKA\KWD\AWNEL
						KGTSKGDAVEA\YLNQV\EELK
	İ			i	ŀ	KKYGDMERLDLVTVPCVYPKP
	i					ETIPFFFSNTVDGWEFRENNPV
						KPATQRLLTHTGS
23248	53616	Α	23382	380	1041	EVTINSGFIFIMLFMEHPEHYTI
		l				MHKWILTWILPTLLYRSCFHIIC
	i					LVGTISLACND\MTPEQMATNV
İ		1				NCSSPERHTRSYDYMEGGDIRV
	l					RRLFCRTQWYLRIDKRGKVKG
	l	1				TQEMKNNYNIMEIRTVAVGIVA
		l			i	IKGVESEF\YLAMNKEGKL\YA
	ŀ	l			1	KKECN\ED\CNFKELILENHYNT
		l				YASAKL/WTHNGGEMFVALNO
		l				KGIPVRGKKTKKEOKTAHFLP
		l				MAIT
23249	53617	Α	23383	127	549	SSPKMAATMKKAAAEDVNVTF
	1	1				EDQ\QKINKFARNTSRITELKEEI
		l				EVKK\KQLQNL\EDACDDIMLA
	1					DDD\CL\MIPY\OLGDVFISHSOE
		1				ETQEMLEEAKKNLQEEIDALES
						RVESIORVLADLKVOLYAKF\G
						SNINLEADES
		1_				SNINLEADES

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SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23250	53618	Α	23384	1	2050	MEYESMVISFDDRVMEKTEYG
		1				PIFTVFAMGNRMTFVTEEEGIN
				•		VFLKSKKVDFELAVQNIVYRT/
						SIPKNVFLALHEKLYIMLKGKM
						GTVNLHQFTGQLTEELHEQLEN
						LGTHGTMDLNNLVRHLLYPVT
						VNMLFNKSLFSTNKKKIKEFHQ
		1				YFQVYDEDFEYGSQLPECLLRN
						WSKSKKWFLELFEKNIPDIKAC
					1	KSAKDNSMDSDEDPVNLQSRS
						ALVCEMLHVQRSSSDADAIGPI
						ADHSLSNRWEVILADEHGIIAD
						DMISRYALSEKSQVELSTQVIKS
		1				ASSKSVNKSDIDTSVFLNWYNF
	1	1				EKKMIRAYATVIDGPEYFWCQI
	1	ı				ADTEKLQCLEVEVQTAGEQVA
		1			ł	DRRNCIPCPYIGDPCIVRYREDG
		1				HYYRALITNICEDYLVSVRLVD
		ı		į.		FGNIEDCVDPKALWAIPSELLS
		1				VPMQAFPCCLSGFNISEGLCSQ
		1				EGNDYFYEIITEDVLEITILEIRR
	l	1				DVCDIPLAIVDLKSKGKSINEK
ŀ		1				MEKYSKTGIKSALPYENIDSEIK
		1				QTLGSYNLDVGLKKLSNKAVQ
l		1		i		NKIYMEQQTDELAEITEKDVNI
	!			1		GTKPSNFRDPKTDNICEGFENP
l	ļ	1				CKDKIDTEELEGVSQKAQESMO
l						TEDMRKSSCVESFDDQRRMSL
l		1				HLHGADCDPKTQNEMNICEEE
		ł				VEYKNRDAISALMP\FSLRKKA VMEASTIMVYQIISQLNYRTPTI
23251	53619	A	23385	3	602	TRPPTRPGLSPDPTSPTEADRGV
23231	33019	^	23363	ľ	002	CERPHRCFLLPQITHPSSGRRCL
l						CLLSPTSFDFSLLYESOSDRAIF
		1				GVGTEPDQEPPPPYHEQVPVSV
		1				YHPT\P\SQTWVATQLTEEEQIR
		1				AQRIGLIQHLPKGVY\DPWRDG
		1	١.			SEKKIRECVICMMDFVYGDPIR
				_		FLPCVHEWYHLDCIDDWLMRS
		1				FTCPSCMEPVDAALLSSYETN
23252	53620	la-	23386	33	512	TAARCQGGPTHVPAFVGTACP
23232	153020	1	23300	155	l	TRIPGHAPAROCREYYY*YYYE
		1		1		F/WKEPRRCHALLLPPWAGLEV
		1			1	GMPPSHTQLGMHPAPPPPHLEV
		1		1	1	KAQG\PSNRQRTDGCRWCSGR
		1				ASALGGGRODSRLRERGGGYD
		1		1	1	H/HCPICRSLKIOLKCHPASGPO
		1		1	l	TVGLGRLASVS
23253	53621	A	23387	1	1731	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:		Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	tocation of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
23255	53623	Α	23389	ī	1986	MGIGEGPTTSEGCRESVEYSHQ
				ŀ		KSELPLSGDSQVPGPIGCNCDP
						QGSVSSQCDAAGQCQCKAQVE
						GLTCSHCRPHHFHLSASNPDGC
						LPCFCMGITQQCASSAYTRHLIS
						THFAP/GDFQ/GFA/LVNPQRNS
						RLTGEFTVEPVPEGAOLSFGNF
						AOLGHESFYWOLPETYOGDKG
						EAYFARMRRAHONRTSLSEEO
						LRAAVTAGRIPEPPEGRDWAQR
						ASOFSLSYEGFSLLPGSLYYWO
						LPRAFLGDKVAAYGGKLRYTL
						SYTAGPOGSPLSDPDVQITGNNI
						MLVASQPALQGPERRSYEIMFR
i			İ	l		EEFWRRPDGQPATREHLLMAL
1	ŀ		i	l		ADLDELLIRATESSVPLAASISA
ĺ		l		i		VSLEVAQPGPSNRPRALEVEEC
i		1		i		RCPPGYIGLSCODCAPGYTRTG
ł	ŀ	l		ĺ		SGLYLGHCELCECNGHSDLCHP
1						ETGACSQCQHNAAGEFCELCA
		l				PGYYGDATAGTPEDCOPCACP
		l				LTNPENMFSRTCESLGAGGYRC
		ı	ļ			TACEPGYTGQYCEQCGPGYVG
ŀ		l				NPSVQGGQCLPESSELHFPSVQ
ŀ		l				PSDAGVYICTCRNLHOSNTSRA
		Į			1	ELLVTEAPSKPITVTVEEQRSQS
					I	VRPGADVTFICTAKSKSPAYTL
		l			i	VWTRLHNGKLPTRAMDFNGIL
						TIRNVQLSDAGTYVCTGSNMF
1						AMDOGTATLHVQGSAEAGERS
		l				MYSISVLLKYTS
23256	53624	A	23390	3	13319	PASGRIRAARSGRRGASERESG
23230	33024	ľ`	23370	Ĭ	13317	AGRAMGWRAPGALLLALLLH
1						GRLLAVTHGLRAYDGLSLPEDI
1		1			1	ETVTASQMRWTHSYLSDDEYM
		1			1	LADSISGDDLGSGDLGSGDFQM
					1	VYFRALVNFTRSIEYSPOLEDA
ŀ					İ	GSREFREVSEAVVDTLESEYLKI
						PGDQVVSVVFIKELDGWVFVE
1						LDVGSEGNADGAOIOEMLLRVI
	1		1			SSGSVASYVTSPQGFQFRRLGT
	1		1			VPOFPRACTEAEFACHSYNECV
	1	1	1		1	ALEYRCDRRPDCRDMS
	<u> </u>		L	L		MLE I KUDKKPDUKDIVIS

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23257	53625	A	23391	41	1323	HYLVFFLPSLRRYTVSLSPNSTS YQLLVPGPSY/SALPISSAASLY AGA/GGGGSSRMSVSHSTSLRDG ISSGGLAVTGMARGLGGMGGIQ NKKETTQSLNDSLAVRVRSLET ENRKLESKIREHLEKKGPQVRD WSHYFKTIEDLRAQIFANTVDN VRIFQIDNARLAADDFRVKYE TELAMRQSVESDIVGLHKVIDD TNVINGCSFFMKKNHGEEVKC LQAQIASSGLTMEVDAPKSQDL AKTTADISTRYDELARKNREEL DKYWSQPIEESTTEVTMQSAKV RAAEMTFMELRRTVQSLEIDLD LMRNPKASLENSLREVEACYA LQMEQL/NLDP/GILLHLESELTQ
						LEAEIATYCLLLEDGKDFNLGD TLDSSNPMQTIQKTTTSRIVDG KVVSETNDTKVWRH
23258	53626	Α	23392	1	77 I	
23259	53627	A	23393	732	1656	SCSWTSPAAWPPNASLHSCLKR RNOPHLLWUWRFLDFSSMA SKCQSPLLPQKKKPRPLPALGP EKTSAAGLPKKGEKEQQETIE HIHEVQNEMDKINEQAKRSEL ISKIPDFWYTFVNHPEVSALPG EEDEEALHYLTRVEMTDFDDIK SGYRIDFYFDENPYFENNILAKE FHLNESGDPSSKSTEIKWKYGK DLMKRSSGOPINKAHRKRQHEE PESFTWFTDHSDVGADELGEV IKDIWPNLQYVLVPDMONEGE GEGEDDDDDEEEGGLEDIDEEG DEDGGEGEDEGERGEEDBGE DEDGGEGEDBGERGEEDBGE DEDGGEGEDBGERGEEDBGE
23260	53628	A	23394	3	923	ARRLYWWFWDFPNSMAPKRGS PLPPOKKKPRPPPALGBEETS AGLPKKGEKEQGAIEHIDEVQ NEIDRLNEQASEELIKVECKYN KLRQPFFCKRSKLISAGIPNFWV TTFVNIHPQVS\ALLGEEDEEAL\ HYLITRVEVTEFEDIKSGYRIDF FYDENPYFSKVLISKEFILINES GDPSSKSTEIKWKSGKDLTKRS SQTQNKASRKRQHEEPESFFTW FTDHSDAGADELGEVIKDDIWP NPLQYYLVPDMDDEEGGEED DDDDEEEGLEDIDEEGDEED EDEDDDGEEEGLEDIDEEGDEED

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	bod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23261	53629	Α	23395	732	1656	SCSWTSPAA WPPNASLHSCLKI
				1		RNQDHLLLWDWRFLDFPSSMA
	ŀ		1	[SKCOSPLLPOKKKPRPLPALGP
	l					EKTSASAGLPKKGEKEOOETIE
				1		HIHEVONEMDRLNEOAK\RSEL
				İ		ISKIPDFWVTIFVNHPEVSALPG
				l		EEDEEALHYLTRVEMTDFDDIK
	İ	ĺ		l		SGYRIDFYFDENPYFENNILAKI
						FHLNESGDPSSKSTEIKWKYGK
						DLMKRSSQTQNKAHRKRQHE
						PESFFTWFTDHSDVGADELGEV
				1		IKDIWPNPLQYYLVPDMDNEE
		1				EGEEDDDDDEEEEGLEDIDEEG
		1				DEDEGEEDEDEGEKGEEDEGE
23262	53630	A	23396	3	924	ARRLVVWFWDFPNSMAPKROS
23202	33030	l^	23390	ľ	724	PLPPQKKKPRPPPALGPEETSAS
						AGLPKKGEKEQQ\EAIEHIDEVO
l						NEIDRLNEQASEEILKVEQKYN
		1				KL\RQPFFQKRSK\LSAGIPNFW
					1	VTTFVN\HPQVS\ALLGEEDEEA
ļ						L\HYL\TRVEVTEFEDIKSGYRID
ļ						FYFDENPYFENKVLSKEFHLNE
				l		SGDPSSKSTEIKWKSGKDLTKR
ļ		1				SSQTQNKASRKRQHEEPESFFT
		l			i	WFTDHSDAGADELGEVIKDDI
ŀ						WPNPLQYYLVPDMDDEEGEGE
	1		}			EDDDDDEEEEGLEDIDEEGDED
						EGEEDEDDDEGEEGEEDEGED
23263	53631	A	23397	1	1393	MGKQKGGGPQAWEAEGLGRE
23203	33031	^	23351	l'	1555	WMRMPQEGKGSQRKSTQKQT
1				l		KREIDA YIVQAKERSYETVLSF
						GKRGLNIAASPAVQAATKVLW
				1		APSPPAAPIPPLRPFGLIQLLCQC
						AQRWQPGSREVKGRLCQARLA
				1		ETSSIAGNLRVQSYERPVDPGV
		ı			1	TMDPLLPGRTIVFAAGDHTLW
		ı				RAPVSSCAQRSGVWSLDNMGV
						NSGSVPFHGVRIQASQLISLSFN
		1				FSFCEADLKTPLHGFLILPAGPK
		1				EEVTVKKEKRERDRDRQREGH
ļ		l				GRGRGRPEVIQSHSIFEQG\PAE
		i				MMKKKGNWDKTVDVSDMGPS
l		1	1		1	HIINIKKEKRETDEETKQILRML
l		1			1	EKDDVDHHALSEHLVTTATFSI
		1			1	GSTGLVVYDYQQVECGGHVRA
		1			1	DLKTQDLYSHAQFGDNNYPGG
					1	VDWYRAGGLQDSDTEDECWS
		1			1	DTEAVPRAPARPREKPLIRSQSL
1		1	1			RVVKRKPPVREGTSRSLKVRTR
		ļ.	22200		504	KKTVPSDVDS
23264	53632	A	23398	297	584	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23265	53633	A	23399	2	149	CMKFGAVTPIGGPPLGDQSPVL
						LLFAL*KRSTYNLGSSDPPAQGI
		1				SHOF
23266	53634	l _A	23400	1	170	PAGIRHEERSTYDLWSSDRPGQ
						GTSHOF*IIRFOIPWNSGPRLSE*
		1				LLPRSSRLSS
23267	53635	Α	23401	I	1077	
23268	53636	В	23402	289	1359	
23269	53637	Α	23403	3	2233	
23270	53638	Α	23404	1	1491	
23271	53639	В	23405	76	511	
23272	53640	Α	23406	1	2706	
23273	53641	Α	23407	273	862	
23274	53642	Α	23408	1	1080	
23275	53643	Α	23409	134	675	MHYFHSSVQIGLPLTRRSQVLIL
		1				PQPLLPNPIILLSPPLLTPVPAYS
l		1				FIL/PTSPPPPARQFPLKKVAEAK
		1		1		GIVKQLKTDTARSPRKPTGPSQ
				1		TLWVTLTVEDLKHAFFTIPLHP
		1		İ		SSQPLFAFTWTDPDTHQAQQIT
		1		1		RAVLPQVFTDSPHYFSQAQISSS
						SVTYLGIILIKTHVLSLLIMSG
23276	53644	Α	23410	1	895	MEVHKHPFNTLALRHESGCEL
				•		VWAEKHRNQIFPFPKSTASPKQ
	ļ		1			WDTGRHNSDAKQQEPGGREER
		1				NTMEMETKETHFIRGPKTLASV
		1				TDWEGSLPLVFNHCRGTSLIIHP
		1				RFKGVRPRRDACLGPSPLAASP
		1	1			AFLGKGQHALKGLKPVITRLLQ
						HGLLKPINSPFPS\PILPVLKPDK
			i			AYKLVQNLRLINQIVLPIHPVVP
		1				NPYTLLSSIPPSTTHYSVLDLKH
		1				AFFTIPLHPSSQPLFAFT*TDPDT
		1	İ			H*AQQITWAVLPQGFTRQPPLL
						QLSPNFILICYLSRHNSHKNTRA
						LPADRV
23277	53645	Α	23411	1	930	MIQEHADKQVQRLQGVLLGSIP
						TAASRARLHPGEINNHVAHTEP
		1				VWWSLHMDAYEIWCRGSDRR
		ı				TSLGRSIPLPPALCSLRRIHLRPQ
		1				VLRSTSPRNISPISNPGHLSDYTP
1		1				TFQGCQTMQGRLPWSFTLSGK
	1	1	1			SRFSGEGATQRQYPIPQQALKG
1			1			LMPAITRLLQHGLLKPINSPYHS
			l		1	PILPVLKPDKPYKLVQDLRLINQ
		1	l			IVLPIHPMVPNPYTLLSSIPASTT
1		1	1			HYSV\LYVKHAFFTIPLHPSSQP
						LSTFTWTDPDTHQAQQIT*AVL
		1	l	1		PQVFTDSPHYFSQAQISSSSVTY
		1		1		LGIILHENTRALPADHV

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop eodon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23278	53646	Α	23412	299	1028	RLGSFSSNHPYLGQRQYPIPQQ
		1				ALKGLKPVITCLLQHVLLKPVN
		1		İ		SPYNSPILPVQKLDMPYRLVOD
						LCLINQIVLPIHAMVPNPYTLLT
		1				SIPPSTTHYSVLDLKPAFFTIPLR
		l				LSFQPLFTFSWTDPDTHQAQQI
						TRAVLPQSFTDSPHYFSQAQIS*
		1				RLGSFSSNHPYLGQRQYPIPQQ
		1				ALKGLKPVITCLLQHVLLKPVN
		1				SPYNSPILPVQKLDMPYRLVQD
		l				LCLINQIVLPIHAMVPNPYTLLT
		l				SIPPSTTHYSVLDLKPAFFTIPLR
						LSFQPLFTFSWTDPDTHQAQQI
		l				TRAVLPQSFTDSPHYFSQAQISS
		1				SSVTYLGIPGFAILTKPLYKLTK
		l				GNLADPIDPKSFPHSSFRSLKTA
					1	LETAPILALPVSSQPFSLPTAEV
						WAAQSEFLHKNRDRTE
23279	53647	Α	23413	157	202	RSECARWMEK*CPWL
23280	53648	Α	23414	184	482	EGVSVFLLLSPYQGQPQVSRRC
						S/RQVSQRCPHRGRGPRCPQVS
		ļ				RSHGPSWVGHPGMPLYGS\EQF
						YCGAPGLAGPPNGSRGESLKIS
						VWIFPCRDFGNFN
23281	53649	В	23415	13	862	
23282	53650	A	23416	1	11342	MATRWFYYGHPPRDARVPRIV
	1	1				AEAKEKNLTPGKIIEVDVTIEAQ
		1		ľ		IRQEFLDAERDTGGFVKFFRITK
		1				KKLQEFFLGEEEEQVPHVSPVE
		1				LGAKPTAFNAFIANAVGKTLAV
	1	1				QHASLKAPEPEMTVTHERFYG
		l		l		NYAVWITDVDPRNHRIEVIHVI
						REISGRTSEYVNNLLAQPTPFK
		1		ļ		VFDWLTETQAKEFVRRLNNTQ GVISVYNKLEDGIPVKVAVKAT
						EPTNKEKLEAASTKLKAMGAV
						ATDAYEFHTDLNENE
23283	53651	A	23417	7597	7883	LSSLAWVWQWNHRISVLFHSN
23263	33031	^	23417	1391	/883	SVRTNLVPQFNQLVITESNSCPE
į						FFNOTSAFFKT*VESGCM/CLOD
	ŀ	1				*GIEWFRLRLDVYNSRLVFLTG
		1				ATDYCTG
23284	53652	A	23418	3	1573	
23285	53653	A	23419	6692	6793	GASVAPSIGS*STGNAIPSCLFQ
		Γ.		1		QNVIVPIPQS
23286	53654	Α	23420	1	17727	
23287	53655	Α	23421	6687	6785	VNTGNRTWWEPLFRK*KKTQR
		L				EDTTEKKTKGR
23288	53656	Α	23422	1	2856	
23289	53657	Α	23423	1	7194	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23290	53658	Α	23424	7460	7561	GASVAPSIGS*STGNAIPSCLFQ
						QNVIVPIPQS
23291	53659	A	23425	1468	1857	
23292	53660	A	23426	1	1985	MEPVGIDIIAISWIGAPEYPDNV
		1		1		KRAINAKVEATQKTLQREQEIQ
		1				QRVAEANMEREQAKGVADAIR
		1		1		LRGEALTENPNVMQLEAINKW
		1		i		NGLVPFITSSPGMGKSAIVRQIA
		1				EDFGLKLIDHRLSTSAPEDLSGL
		1				PRFKENGRAEFAPFDELFPLEG
		1				DAVPEGFNGWLLFLDEFNSAR
		1				KEIQAAAYKLVLDRMTGQKKL
		1		Į.		HPYVAIVCAGNKATDRAITNNL
1		1				STAMQSRLIHLEMETDFDVFME
		1				DVAIPNKWDERVIAFLNANPNK
1		1		l		LNDFEPDHQEKTFCCSRTWEFV
		İ				NKIVSILPPGPINSEMTLLLAGTI
		1				TSGVATSFVQFTQVYSNMVSL
		1		1	1	NEILDNPKKARMPEDNNLLWA
				l		VVTSLINNTDEDNHSKIFDYVE
		1				RMPFTFKVLYYRSRKTDMSMT
		1		1		DAQLTREYDRCQAKAFMGKTA
		1				AFFGSLLCSLKFRWVKDGCDT
		1				AQTDGEHLEWNPDWFESLLPE
						SRVTVLMHELWHVGLLHSVR
		1				MGSRDPEVWNYACDIYINNQLI
İ		1				QDGYSFIGIENCWKDPKYAGW
l				ŀ		AEEQIYHDLMSKHPRPPKASGA
		1				FGTGSGGDMKPSTSKASLKLSG
		1		1		GSLPGVLPGKMEEVITQFLKPV
i		1				VPWQVLLERFFNDLQETYYSW
		1				QRPNRRYPDMYLPSSMDDDCR
		1				LEHLAYFL*NEADLTGLLVLK*
						TDDGQSVSGQNLLQSDEQLQIY
23293	53661	Α	23427	2365	2784	
23294	53662	Α	23428	5789	5980	DQNMRFDNTFETYYWKPTIIVII
		1		i	i	IRKLLMNDSSKDTLK*LRCWK
						NWLQILKKHTIHFIMPLQL
23295	53663	Α	23429	I	6819	
23296	53664	Α	23430	376	2052	SKVCGNDGFADDLKHLLIFTGA
		1				QSGEDIVPFQLEDGHSLGEVVL
		1	l		1	LHGGSGVESGQR/CGRTSAGSC
		ı				Y*SPGKICCCNLLVTKTEGFFAP
						VPIQLENTNQVTNCSDNW
23297	53665	Α	23431	2	2189	
23298	53666	Α	23432	2	1583	
23299	53667	Α	23433	2142	2242	GASVAPSIGS*STGNAIPSCLFQ
		1_				QNVIVPIPQS
23300	53668	Α	23434	3	1188	
23301	53669	Α	23435	470	862	
23302	53670	A	23436	1	1352	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	deletion, \=possible nucleotide insertion)
23303	53671	A	23437	1837	1955	SKCSIGLYQRYPYSNGC*NSGCI CRVY*CCREHRGYVRN
23304	53672	A	23438	1447	1565	SKCSIGLYQMYPYSNGC*NSGC ICRVY*CCREHRGYVRN
23305	53673	A	23439	773	823	
23306	53674	Α	23440	3	1529	
23307	53675	Α	23441	1	1794	
23308	53676	A	23442	236	427	DQNMRFDNTFETYYWKPTIIVII IRKLLMNDSSKDTLK*LRCWK NWLQILKKHTIHFIMPLQL
23309	53677	A	23443	8510	8611	GASVAPSIGS*STGNAIPSCLFQ QNVIVPIPQS
23310	53678	A	23444	516	669	ATSELKRKASLGLMEPDVSRK* ARCSSLPSSSIGDGRYISGYLRL GRCHE
23311	53679	A	23445	410	593	FLNLIQPQGKRKLLLKRCRSLLP KN*CQHSKQHKLSISSVVILNIT NS*PNKVKLMPILAM
23312	53680	В	23446	1	2286	
23313	53681	Α	23447	1418	1723	CNTNIIFNCHFPTKLGCICKAMK VVSPLIHNSIVILLNQITIDFFSTL AQICTFICLTYSKQTTTRIASFK KVIEEFLSLAPIEKVELKEHS*YI SISEKV
23314	53682	В	23448	1	6167	
23315	53683	Α	23449	1	3199	
23316	53684	Α	23450	1	5733	
23317	53685	Α	23451	709	1099	
23318	53686	A	23452	5039	5230	DQNMRFDNTFETYYWKPTIIVII IRKLLMNDSSKDTLK*LRCWK NWLQILKKHTIHFIMPLQL
23319	53687	Α	23453	664	993	
23320	53688	A	23454		19859	MASIDSIAVCNPRQVYKFVEHC IRSGLVPFITSSPGMGKSAIVRQI AEDFGLKLIDHRLSTSAPEDLSG LPRFKENGRAEFAPFDELFPLEG DAVPEGFNGWLLFLDEFNSAR KEIQAAAYKLVLDRMTGQKKL HPVVAIVCAGNKATDRAITNNI STAMQSRLIHLEMETDFDVFME DVAIPNKWDERVIAFLNANPNK LNDFEPDHQEKTFCCSRTWEFV NKIVSILPPGPINSEMTLLLAGTI TSGVATSFV
23321	53689	В	23455	74	240	
23322	53690	Α	23456	143	384	
23323	53691	Α	23457	1	3267	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide scquence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \-possible nucleotide insertion)
23324	53692	A	23458	2	550	KRSSEHIQRIDRDVSGTLRKHIF FRDRYGTKQRELLHILLAYEEY NPEVGYCRDLSHIAALFLLYLP EEDAFWALVQLLASERHSLQG FHSPNGGTVQGLQDQGEHVVA TSQPKTMGHQDKKDLCGQCSP LGCLIRILIDGISLGLTLR'LWDV YLVEGEQAFMPITRIAFKVQQK RLTKYSR
23325	53693	Ā	23459	72	630	
23326	53694	A	23460	I	2835	MDALMLALGALMILAGLOD KSAQVSDPDGAEPGPGRATGT DPLVQIIIWKLGISEALSIPGGRL YRDLWSLTPVCLHIPGIAHHGP FTLGRMDVVEVAGSWWAQER EDIIMKVEKGHRAGLPEDKGPK SFGSYNNNVDHLGIVHETELPP LTAREVKQIRREISRKSKWVKM LGEWDTYKNSKKLIDRAYQGIP MNIRGPMWSVLLNIEEIKLKNP GRYQVRSARAQPTGQAVSGAQ VSSWRERQDHPGELGVKI

3075

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	tocation of first	codon for fast amino acid	*=Stop eodon, /=possible nucleotide
	sequence	ı	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1				sequence		
23327	53695	A	23461	li .	832	MGWRARWLMPVISALWEAED
		1		1		PIPPELTRLLQQSQDPLLMGLFP
		l				TNPKEKTQEEPPGQSRAPVLTV
		1				VSKFKVLAGLLHLGNIQFAASE
		1				DEAQPCQPMDDAKCEGQGVG
ĺ		l				WAGPVRKASLEQLLQVLHSTT
		l				PHYIRCIKPNSOGOAOTFLOEE
						AGAGRRRORRROKOORRRRRO
		l				LGGGGDGGNGSVGDDGATRGS
		1				RGCALEKECVATPRRGLLFSRP
		l				HKLAFAVGGSVTSVGPKPRAP
		l				MDVVEVAGSWWAQEREDIIM
		l				KYEKGHRAGLPEDKGPKPFRS
1		l				YNNNVDHLGIVQDFADFGTTIK
1		1				QDFRLLGQTSVDRLLQLSQGQ
ļ		l				AVKGNQLLPVSLVKRKTTLAP
1				1		NTQTASPRALADSLMQLARQV
		l		1		SRLESGHFYDAHNPFLPPPSMS
		1				KQAKKTGVPSEGLCRPGGEGQ
		1		1		SAVARLTGEKKNHPGAQYANR
		l				LSPRVGRFINAAGTTGFPTGKR
1	l	l				AVSATQLIGLQLILLGEVLQKV
		1				LTASYPAQSSVPLTSGPVRQIM
1		1		i		KEKGKRSSEHIQRIDRDISGTLR
i		1				KHMFFRDRYGTKQRELLHILLA
		1		ļ		YEEYNPEVGYCMDLSHIAALFL
		1				LYLPEEDAFWALVQLLASERHS
	ļ.	l				LQAARAPAAIGAHERADQAQIS
		1				LGLTLRLWDVYLVEGEQALMP
		l				ITRIAFKVQQKRLTKTSRCGPW
1		l				ARFCNRFVDTWARDEDTVLKH
		-				LRASMKKLTRKQGDLPPPAKPE
23328	53696	A	23462	3	497	SSECLTMAWIPLLIPLLILCTVS
						VASYELTQPASVSVSPGQTASIT
1						CSGDVLAKKKYARWFQQKPG
						QAPVLVIYKDSERPSGIPERFSG
		1				SSSGTTVTLTISGAQVEDEADY
		1				YCYSAADNNGWVFGGG\TKLT\
		1				VLSQPKAAP\SVTLFP\PSSEELQ
23329	52607	١.	22462	1	393	ANKAHTGVSQ
23329	53697	A	23463]'	393	FRSGSELVKAGLRAFFENAAED
		1				LEKTSENLKLGKFTHSRTQIKG
		1				VSQNINYTTVALLPILTSIFEHV
		1				TQHQFGMDLLLGDVQISCYHIL CSLYSLGTGKNIYVE\ILGMPDT
		1				
23330	53698	Α	23464	319	508	VEDMCPDIPQLECLMKEIN
23331	53699	C	23465	94	222	,
23332	53700	A	23466	1	636	
23333	53701	Â	23467	1	483	
233334	53702	Â	23468	1	1797	
~5554	100/02	10	22700	ı·	1.121	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23335	53703	Α	23469	l	4275	
23336	53704	Α	23470	l	1848	
23337	53705	Α	23471	122	14928	ARSWRRRWRTRTKRRRKSAA
		1				AVSARRAAAGGSRGAGGWGT
						ADASGAMAEGGEGGEDEIQFL
1						RTEDEVVLQCIATIHKEQRKFC
						LAAEGLGNRLCFLEPTSEAKYIF
						PDLCVCNFVLEQSLSVRALQEM
						LANTGENGGEGAAQGGGHRTL
		1				LYGHAVLLRHSFSGMYLTCLTT
						SRSQTDKLAFDVGLREHATGE ACWWTIHPASKQRSEGEKVRIG
		l		l		DDLILVSVSSERYLHLSVSNGNI
	İ	1				OVDASFMOTLWNVHPTC
23338	53706	A	23472	134	1063	RPPGLTRSPSSPAPARSPTRRQS
23330	33 /00	^	23472	134	1003	TRRRSGATTSCSQRLWATWNG
						CD/VCLNQSLREIPTDDKGFTAI
l		l				HFAAQWGKLACLQVLVEEYKF
						PVDLLTNNSQTPLHLVIHRDNT
						TVALPCIYYLLEKGADLNAOTO
	1					NGSTPLHLAARDGLLDCVKVL
		1				VOSGANVHAQDAMGYKPIDFC
		l				KIWNHRACARFLKDAMWKKD
						KKDFAREMTKMKMFKSQLTL
		1				MEHNYLIEYQKEHKILREAAIR
		1				KWLHGKLHPGHSLVSNTKQAR
						ATALSKTPEQRESQRSRSFHPSV
	1					DARLQCIPQPTEMPKPIYRKPTV
						KRPHNVEC
23339	53707	Α	23473	1	1280	MAGLGFWGHPAGPLLLLLLLV
					1	LPPRALPEGPLVFVALVFRHGD
1					1	RAPLASYPMDPHKEVASTLWP
ļ						RGLGQLTTEGVRQQLELGRFLR
		1				SRYEAFLSPEYRREEVYIRSTDF
		1				DRTLESAQANLAGLFPEAAPGS PEARWRPIPVHTVPVAEDKLLR
		l				FPMRSCPRYHELLREATEAAEY
		l			j	QEALEGWTGFLSRLENFTGLSL
					1	VGEPLRRAWKVLDTLMCQQA
	1					HGLPLPA WASPD VLRTLAQISA
					1	LDIGAHVGPPRAAEKAQLTGGI
	1	1				LLNAILANFSRVORLGLPLKMV
1	1	1			1	MYSAHDSTLLALQGALGLYDG
	1				1	HTPPYAACLGFEFRKHLGNPAK
		1			1	DGGNVTVSLFYRNDSAHLPLPL
		1			1	SLPGCPAP\CPLGRFYQLTAPAR
	1	1			1	PPAHGVTCHGPYEAAIPPAPVV
	1	1			1	PLLAGAVAVLVALSLGLGLLA
1	1	1			1	WRPGCLRALGGPV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23340	53708	A	23474	133	672	AVGWESGK WDSGRGQGGGQE
						OMWSCOSDMLYPPDOVGAWG
	l					LPGPTGPKGDAGSRGPMGMRG
ŀ	i i	l				PPGECPQCCPTAPLAIPSTRWAL
l		l				PCSWPPGPPGPPGPPAPVGPPHA
		1				\GSPSMVSPPGIPAGGGGGGVAI
	Į.					STVPATIKTRROEVSPELPRP/PG
		1				PAPPPGPDVRPWLLRPPGPMGK
						LSPGQG
23341	53709	Α	23475	135	677	AVGWESGKWDSGWGQGGGQE
	i	1				QMWSCQSDMLYPPDQVGAWG
						LPGPTGPKGDAGSRGPMGMRG
		1				PPGECPQCCPTAPLAIPSTRWAL
						PCSGPPGPPGPPGPPAPVGPPHA
		ı		ľ		\GSPSMVSPPGIPAGGGGGGVAI
ŀ		1				STVPATKKTRRQEVSPELPRP/P
ļ						GPAPPPGPDVRPWLLGPPGPMG
				1		KLSPGQG
23342	53710	Α	23476	220	745	TVIQRTGLHSVGPGDPQQGLRE
l		1				AKLGTMVRLVLLPTPRRTTEVR
						TGPERELSMVLEAILGSLLETAC
	İ	1		1		DHRQTPPSSKPQA/GSSRLREGT
						SAGQPCS/LRLRLGPRAGGRRC
	Į.	1				CRDIERSSSSSVCGRKCKRKKR
		l				KRGMCQSRHKHGCRRPQEGPR
						CNDGRALCSPRPQTTPGPLSTA
23343	53711	A	23477	1	2433	
23344	53712	Α	23478	3	803	FPGRRFRVRACVAGGSHCLFNS
	1			ļ		CCCCTSASLPARERHEWPGPAA
	1	1				AAAAAAAARRSQTQREGLFA
		1			l	GWGGGFAMSDDDSRASTSSSSS
						SSSNQQTEKETNTPKKKESKVS
		1				MSKNSKLL\SPRAKGIQKEL\AD
		1				ITLDPPPNCSAGPKGDNIYEWR
		1				STILGPPGSVYEGGVFFLDITFTP
						EYPFKPPKVTFRTRIYHCNINSQ
			1			\GVICLDILKDNWSPALTISKVL
			1		1	LSICSLLTDCNPADPLVGSIATQ
				1		YMTNRAEHDRMARQWTKRYA
	1	1				T

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1				sequence		
23345	53713	A	23479	372	1839	GEAGGTEAEPENGAGGPRVPG
23343	33713	^	23419	1772	1037	WWGWSSTSSPRGCWDSGRSAA
1						AIMDFLWDKRTGLAARKMPHP
						RRYHSSERGSRGSYCEHYRS/R
ı						K/HKQRRSRSWSSSSDRTRRRR
i						REDSYHVRRRCSRTFSRSSSOH
						SSRKAKSVEDDTEGHLIYHVGD
1						WLOERYEIVSTLGKGTFGRVV
			[OCVDHRRRGARVALKIIKNVE
1		l				KYKEAARLEIKVLEKINEKDPG
						KNLCVOMFDWFDYHGHMCISL
1		l				ELLGLSTFDFLKDNNHLPYPIH
1		l				OVHHMASOLCOAVKFLHDNK
1		1				LTHTDLKPENILFVNSDYELTY
1		l				NLEKKRHERSVKSTAVRVGDF
1						GSATFDHEHHSTIVSTRHYRAP
1		l		i		EVILELGWSQPCDVWSIGCIIFE
1						YYVGFTLFOTHDNROHLATME
1		l				RILGPIPSRMIRKTRKOKYFYRG
						RLDWDENTSAGRYVRENCKPL
						RQYLTSEAEEDHQLFDLIESML
		ŀ				EYEPAQRLTLGEALQHPFFSRL
						WAEPPNKLWDSSQDISP
23346	53714	Α	23480	28	941	QSARPSLKRARSQRGRPLPSRA
		1				LVFLFLFFFFFFNFLPIHTQPSAH
		1	i			KDMTTNAGPLHPYGPQHL\RLD
		1	1			NFVPN\DRPTWAYTGLGLFSVT
ł		1				GGLRSWTTWAVVQGRAAGCPI
İ		1				GDFGRATVPCCWVLQCVGFIH
ļ						LVIEGLVPFSNYERPCFGDQSLL
		1				NLKLW\KEYGQGETARLHPGG
	1	l			ľ	DKFHSVALGTITSFA\WGEPFSL
	i			1		WVVIA\FLLQHPL\RFIL\QLVVS
		1		1		VG\QIYG\DVF\YFLKEHPRRDS
		1		1		KHGKPGPTPLLLPGFSFVFHEM
		1		1		PLWLVPAPGSPLCLDA\VKAPS
	1			l		THAPEHAWNAKGPQKPKSKKN

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
23347	53715	A	23481	1	1386	MWRNLKLPRDLLNGFDONADS
23347	33713	ľ`	23107		1.500	DMDNEVOAEVVTDGDEELIGN
					·	WSKGHSCYILAKRLAAFFPCPK
						DLWKFELERDDLGYLAEEISSS
						EAFKSVSYEKAHAWNPLESTC
						RHASLALAVVLORRDWENP\G
						VTOL\NRLAAHPPFASW\RNSEE
						ARTDRPSQQLRSLNGEWRLMR
						YFLLTHLCETLVKVKDAEDOL
			1			GARVCVLEROIFDFLGYOWAPI
1			ĺ			LANFVHILIVILGLEGTIOYRPRY
						ITGILSFPKHQESLSLKFTCLTEA
						ERDTDALLHLENACRESTATOR
						IQGFFLKDKKRKKQQPQQQKQ
						OOWAKVOLGPWLORVOAPSL
						GSFHGVLSLRVHRSKELRFRNC
			ŀ			CLDFRGCMEMPGCPGRGVLQA
		l				ONPRGEPLLGOCRREMCGGSP
						YIVPTGALPSGAVRRRPPSSRPH
			i			NGRPTDSLHCVPGKAADTQHQ
						PLKAARRAAVPCKATGSELPK
23348	53716	Α	23482	3	732	SPLTPTATSGNPSKASSPGGLGS
						LVSTASSLPAGEKNSPRTGPGP
						RRAG\AGIPNSGPDTVEGPYNL
			ł			CGRQGRWGGLIISLASEMKLQT
						LQTSVTTAKATLLELFVPLGGL
						VASLASEVRLQIFPRGGHLQSA
			ŀ			KCGLLLHFSLPTSRSGRRKANR
						PMAVTVLLVLIRDCADLLAVK
						AVIHCYQDQWPLRGSSRPANTP
						PPKASVSRRPPGHRGVRRGSPL
						LQVGARRVGAEAPALLELQSPS
	53717	A	23483	1	234	
23350	53718	Α	23484	1	256	MLNLQGKEWLELEEDAQKAYI
						MGLLDRLEVVSRERRLKVARA
		1				VLYLAQDGEKFCPKRDCTNNS/
						RY/NSI*MIQMMRFGTLELMRL
	53719	Α	23485	1	3333	
	53720	Α	23486	1	2661	
23353	53721	Α	23488	3	433	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon fur last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
22254	62720	<u> </u>	02400	3	990	EKMNTADQARVGPADDGPAPS
23354	53722	A	23489	,	1990	,
1		l			ł	GEEGEGGGEAGGKEPAADAA
1		l				PGPSAAFRLMVTRREPAVKLQ
1		l				YAVSGLEPLAWSEDHRVSVST
1		1				ARSIAVLELICDVHNPGQDLVIH
		l				RTSVPAPLNSCLLKVGSKTEVA
		l				ECKEKFAASKDPTVSQTFMLDR
		l				VFNPEGKALPPMRGFKYTSWSP
		l				MGCDANGRCLLAALTMDNRL
		l				TIQANLNRLQWVQLVDLTEIYG
		1	i			ERLYETSYRLSKNEAPEGNLGD
		1				FAEFQRRHSMQTPVRMEWSGI
		1				CTTQQVKHNNESRDVGSVLLA
		1				VLFENGNIAVWQFQLPFVGKES
					1	ISSCNTIESGITSPSVLFWWEYE
					į.	HNNRKMSGLIVGSAFGPIKILPV
					l	NLKAVKGYFTLRQPVILWKEM
		l				DQLPVHSIKCVPLYHPYQKCSC
						SLVVAARGSYVFWRLLLISKAG
		1			1	LNLHNSHVTGLHSLPIVSMTAD
		l				KQNGTVYTCSSDGKVRQVIPIF
		l				TDVALKFEHQLIKLSDVFGSVR
1		l				THGIAVKPCGAYLAIITTEGMIN
1		l			İ	GLHPVNKNYQVQFVTLKTFEE
		l				AAAQLLESSVQNLFKQVDLIDL
1		l				VRWKILKDKHIPQFLQEALEKK
						IESSGVTYFWRFKLFLLRILYQS
						MOKTPSEALWKPTHEDSKILLV
		1	ľ			
						DSPG\MGNADDEQQEEGTSSKQ
		1				VVKQGLQERSKEGDVEEPTD\D
		l				SLPTTG\DAG\GREPMGRRNSW
		<u> </u>				GNSKGKIRSCWEMRLTREHMK
23355	53723	A	23490	1218	1446	RCTDSQV\RSPTPLHLAAQACSL
		l		ŀ		ETTVCLLCSKADYTLSEKRGW
		l				MPIHFAAFYDNVCIIIALCRKDP
				L		SLLEAEATAE
23356	53724	A	23491	2	632	NMAKTYDYLFKLLLIGDSGVG
		l				KTCVL\FRFSE\DAFNSTFIFTIRI
		l				DFKIRTIELDGKRIKLQIWDTAG
	İ	l				QERFRTITTAYYRGAMGIMLVY
		l				DITNEKSFDNIRNWIRNTEEHAS
		l				ADVEKMILGNKCDVNDKRQVS
	1	1	l			KGS/GENKLALDYGIKFMETSA
	1	1	l	l		KANINVENAFFTLARDIKAKMD
1		1	l	l	l	KKLDGNSP\QGSNQGVKITPDQ
		l		-		OKRSSFFRCVLL
$\overline{}$	L	Ц.				Z.C.CO. I IC VLL

NO: of peptide sequence s	SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
23357 53725 A 23492 I I018 MVFYHVGQAGLELLTSGPHCP DSQAADKAESATLLSRARLGA YCACASLERRWHFPSGRECN MAKTYDVLKKLLUBIOSGVGK TCVLFRFSEDAFNSTFISTIGIDF KIRTILELDGKRIKLQIWDTAGQ ERRTDHNRPTYRGAMGIMIL VYDITHEUSSFDONIRNIWIRNIE EHASADWEKMILGONLCOWND KRQVSKERGEKLALDIYGIKFM ETSAKANINVENAFTLLARDIK AKMDKKLGPARPHGTLLGKGE EFAGVAGASASGASLGTLFKQ TRPCPVNPQASRETHATSFILKGE GKDMMEGNSPQGSNGGVKITP DQKRSFFRCVLL 1392 130	1	sequence	l	09/540,217		of peptide sequence	deletion, \-possible nucleotide insertion)
DSQAADKAESATLLSRARLGA			l		sequence		
DSQAADKAÈSATLLSRARIGA	23357	53725	A	23492	1	1018	MVFYHVGOAGLELLTSGPHCP
VCACASLERRWPHFPGRECN MAKTYDYLFKLLLIGDSGVGK TCVLFRSEDAPNSTFISTIGIDF KIRTIELDGKRIKU,QIWDTAGQ ERFRTDHNRPTYRAMGAMGML VYDITHEKSFIDNIRNWRINE EHASADIVEKIMILGNSTOWND KRQVSKERGEKLALDYGIKFM ETSAKANINYENAFFILARDIK AKMDKKLGPARPHGTLLGKGE EFAGVAGASASGASLGTLFKQ TRPCPVIPQASRETHTASIPLKA GKDNMEGNSPQGSNQGVKITP DQQKRSSFFRCVLL			1				
MAKTYDYLFKLLUIGDSGVGK	ł				İ		
TCVLFRFSEDAFNSTFISTIGIDF KIRTILELDGKRIKLQIWDTAGQ ERRRTDHNRPTYRGAMGIML VYDITHEUKSFIDNIRNWIRNIE EHASADWEKMILGNICDVND KRQVSKERGEKLALDIYGIKFM ETSAKANINVENAFTILARDIK AKMDKKLGPARPHGTILLGKGE EFAGVAGASASGASLGTLFKQ TRPCPVNIPQASREHTASTIPLKG GKDMMEGNSPQGSNQGVKITP DQKRSSFFRCVLL				1			
RIRTIELDGKRIKLQIWDTAGQ			l		l		
			i				
VYDITHEIKSFIDNIRNIWIRNIE			i				
EHASADIVEKIMILGNIKCDVND KRQVSKERGEKLALDYGIKFM ETSAKANINVENAFTLARDIK AKMDKKLJPARPHGTILGKGE EFAGVAGASASGASAGATIPKQ TRPCPVNPQASREHTAFSIPLKG QKDMMEGINSPQGSNQGVKITP DQKRSSFFRCVLL			l				
RRQVSKERGEKLALDIYGIKFM ETSAKANINYENAFTILADIK AKMOKKLGPARPHGTILLGKGE EFAGVAGASASGASLGTLPRQ TRPCPVNPQASRETHATSIPLKAG QKDNMEGNSPQGSNQGVKITP QQKRSSFFRCVLL			l				
ETSAKANINVENAFFILARDIK AKMKKLGPARPHGTILGGE EFAGVAGASASGSASLGTLPKQ TRPCPVNPQASREHTAFSIPLKG QKDNMEGNSPQGSNQGVKITD DQQKRSSFFRCVLL			l			l	
AKMDKKLGPARPHGTILGKGE			l				
EFAGVAGASASGASLGTLPKQ TRPCPVNPQASREHTAFSIPLKG QKDMMEGNSPGQSSNGGVLITP DQQKRSSFFRCVLL			l			İ	
TRPCPVNPQASREHTAFSIPLKG QKDNMEGNSPQGSNQGVKITP QKDNMEGNSPQGSNQGVKITP QQKRSSFFRCVLL			i				
QKDNMEGNSPQGSNQGVKITP			l				
DQQKRSSFFRCVLL			l				
23358 53726			l				
23369 53727 A 23494 I 1392	23358	53726	Ι	23493	1	130	DQQRRSSITRE VEE
23360 53728 A 23495 445 649 YPSLTNSKLLVHHFMELCWD KCVEKPQGIALDSRENCLSSC WDRFIDTILATISFR-QUVOKG QQ VIVAPGKKAQRTQK/APKTKTK KPHLSPCPSP/WRSRSCQPMQQS WSGAQVHIPQPPHTTCITWA VIVAPGKKAQRTQK/APKTKTK KPHLSPCPSP/WRSRSCQPMQQS WSGAQVHIPQPPHTTCITWA QASSSPISSVTCKALPPGQTK KGLVASQPEPVSMGTSSPTPS LAPEHSSRASLEPSSSA*PSLAT* SRAGGSQA QASSSPISSVTCKALPPGQTK GGLVASQPEPVSMGTSSPTPS LAPEHSSRASLEPSSSA*PSLAT* SRAGGSQA QAMHGRSPGVTVRASGLRNEG DGPLGSVPTQARATGAPGHSAT PSPAWRAW/SSSSKAS/WLPDA AGPPTPPWSGFSSPTSW QAMFGSSPTSSW QAMFGSSPTSSWASWSSSKAS/WLPDA AGPPTPPWSGFSSPTSW QAMFGSSPTSSWASWSSSKAS/WLPDA AGPPTPPWSGFSSPTSW QAMFGST QAMFGST QAMFGST QAMF			-		-		
XCVEKPQGIALDSRTENCLSISC VIDRIBITYLAITSRFAQIVQKG QQ QQ QQ QQ QQ QQ QQ					<u> </u>		YPSLTNSKLLVHH\FMELC\WD
VORFIDITALATISRFAQIVQKG QQ Q23361 53729 A 23496 210 397 VIYAPGKKAQRTQK/APKTKTK KPHLSPC'SPIWRSRSCQPMQQS WSGAQVHIPQPPHTCITWA QASSSPISSVICKKALPPGQTK KGLVASGDPEPVSMGTISSPITS LAPPHSSRASLEPSSA*PSLAT* SRAGGSQA S3731 A 23498 780 1022 QAMHGRSPGVTVRASGLRNEG DGPLGSVFTQARATGAPGHSAT PSPAWRAWSSSSKASWLPDA AGPPTPPWSGPSPTSW QAMPPTPWSGPSPTSW QAMPPWSGPSPTSW QAMPPTPWSGPST QAMPPTPWSGPST QAMPPTPWSGPST QAMPPTPWSGPST QAMPPTPWSGPST QAMPPTPWSGPST QAMPPTPWSGPST QAMPPTPWSGPST QAMPPTPWSGPST QAMPPTPWSGPST QAMPPTPWS			ľ		1772	047	
CQ CYPAPGKKAQRTQK/APKTKTK KPHLSPCPSP/WRSRSCQPMQQS WSGAQVHIPQPHITCITWA			l		l	ľ	
23361 53729			l				
KPHLSPCPSP/WRSRSCQPMQOS KPHLSPCPSP/WRSRSCQPMQOS WSGAQVHIPQPHITTCITWA	23361	53729	A	23496	210	397	
WSGAQVHIPQPHTTCITWA			Γ.				
23362 233730			ı				
KGLVASGDFEPVSMGTSSPTPS	23362	53730	A	23497	55	286	
LAPEHSSRASLEPSSSA*PSLAT* SRAGGSQA			l				KGLVASGDPEPVSMGTSSPTPS
SRAGGSQA			l				
23363 53731 A 23498 780 1022 QAMHGRSPGVTVRASGLRNEG DGPLGSVPTQARATGAPGHSAT PSPAWRAW\SSSSKAS\WLPDA AGPPTPPWSGPSPPTSW 23364 53732 A 23499 I 782			ı				
DGPLGSVPTQARATGAPGHSAT PSPAWRAWISSSSKASWLPDA AGPPTPPWSGPSPPTSW 23364 53732 A 23499 I 782	23363	53731	A	23498	780	1022	
PSPAWRAWISSSSKASWLPDA AGPPTPPWSGPSPPTSW 1 782 1 782 1 1 1 1 1 1 1 1 1			ľ				
23364 53732 A 23499 I 782			I	l	l		
23364 53732 A 23499 I 782	1		ı		i		AGPPTPPWSGPSPPTSW
23365 53733 A 23500 275 688	23364	53732	Ā	23499	I	782	
	23365	53733	A	23500	275	688	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ı	09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
1		l		sequence		
		<u></u>				
23366	53734	A	23501	1166	2271	LRCILTLISRWCS*VSTVANLPP
		l				GISKWGANWQRCEMKE*CWSP
1		1			i	AVTWCITCAQRSG/IGDSSPYPG
		l				ATSFNEYVKANLTWRGPVEQH
		1				PLVNYLDHERGTLSNPTARHYL
1		l				PVLYVLRAILWARTPLPIPGRTT
1	1	l	:			VSKVKMKKRILRAMRQKQQVL
i		l				PLLVGLGLDELSMSAPSIPAAK
		l				ARMAOLDSRECRKLLNOAMA
	l	l				CRTSLEVEHLLAQFRMTQQDA
		1		i		PLVTAECITLESDWRSKEEVLK
		1				GMTDNLLLAGRCRYPRKLEAD
	ļ	l				LWAREAVFSTGLGFSFAIPHSK
	ļ	l			1	SEHIEQSTISVARLQAPVRWGD
		İ			1	DEAOFIIMLTLNKHAAGDOHM
		l				RIFSRLARRIMHEEFRNALVNA
		l				ASADAIASLLNMNWTVKEHHE
23367	53735	A	23502	1	2604	ASADAIASEENWINWTVKEITIE
23368	53736	A	23502	306	468	TRHCDRROPRRGHRRAGT/CNO
23300	33730	l^	25505	300	1700	I*KGCCCVRTKQPLSSVLISPNF
i		1				CPCSSFLKNS
23369	53737	A	23504	102	509	AAQQGTMARSLLLPLQILLLSL
23303	33131	<u>۱</u> ^	23304	102	1509	ALETAGEEAQGDKIIDGAPCAR
1	1	1				GSHPWQVALLSGNOLHCGGVL
1		1			1	VNERWYLTAAHCKMNEYTYH
		1				LGSDTL/E/ADRKAQRIKPSKSF
		l				RHPGYFTOTHVNDLMLVKLNS
		l				OARLSSMVK
23370	53738	A	23505		1032	MMQATVGFEDGDAATSQVSPS
23370	23/36	Ι^	23303	1	1032	VNKVHSSIYFLDFFEEYIVIAFKI
		l				
		ı			Į.	LRRGPDISCSQIINGEDCSPHSQP
I		1			l	WQAALVMENE/LFCSGVLVHP
ł		l l			ľ	QWVLSAAPVSRKYNRPLLAND
1		1				LMLIKLDESVSESDTIRSISIASQ
1		l			ł	CPTAGNSCLVSGWGLLANDPT
		l				SR\LPPPLECRNPVPQLLLPPEKP
ł		1				RTAPDTLGSSPVPKSRTGSSGSS
1					İ	FLLPAPPSPFPRPHHGTPPTSCG
1						QDVDVPALAGGSLGSPAGEAL
1		1			i	PRLAAPERLSRHHGKIPSPAPAD
		1	l			LTAILALETAGEEVPFPLTSCA
l		l	1		l	WMSSSSPPGLHEGLQGLTGKFH
1						AVRWHPRLQEKRLQSVPSPDSR
		1				IPGARRPAAR
23371	53739	Α	23506	303	393	
$\overline{}$		_				

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \≈possible nucleotide inscrtion)
				sequence		
23372	53740	A	23507	391	1170	GSFFFFFKYFAYSPMPKNSKVV
23312	33740	Α.	23307	391	1170	KRELDDDVTESVKDLLSNEDA
				i		ADDAFKTSELIVDGQEEKDTDV
						EGK\SEVEDERPAWNSKLQYIL
						AQVGFSVGLGNVWRFPYLCQK
		l				NGGVVEPECEQSSATTYYWYR
			ľ			EALNISSSISESGGLNWKMTICL
		ŀ				LAAWVMVCLAMIKGIQSSGKV
		ŀ				NTYDRVTMSATVLALQEQWQT
						FSPIRSGNGHLTGSGAQRTPCRI
						RRGGSQLRVCDGGKQQWWTV
		_				SQSSARAITNTDQKSVQLQDLIE
23373	53741	Α	23508	45	546	EGGGSRSLWFRELIGQPVQSFIE
						PVAAGGAVALNVQ/RSCGVVS
						AAPACL*SRPHSWRWGGPACW
						RTPGG/LASRSSSSASILMSSSRA
			ļ			SFTRSLSLLSTHKDQGLVCSGSS
		1	l	ĺ		GATGAGSC/LWPPTSHTVKLMF
		1				LYSTVSTLKPMVGMVVTISPSL
1		l				SL*RIVVFPAASRPT/YEDAHFFF
						CQKGP*RGSWSSLQHPGPP
23374	53742	A	23509	1367	2142	VRRPQGCLGKRSLLSRVCALVP
		1				WGIWP/WSLAVGPQYSSLGSQP
		l				ILCASIPGLVPKQLRFCRNYVEI
						MPSVAEGIKIGIQECQHQFRGR
						RWNCTTVHDSLAIFGPVLDKG
						YFGYLETITVPSDSVVSTKAETP
						DKGVCGRELGFRGGWLPELQS
						LDEISMGWALCSRLQIKATRES
		l				AFVHAIASAGVAFAVTRSCAEG
						TAAICGCSSRHQGSPGKGWKW
		1				GGCSEDIEFGGMVSREFADARE
						NRPDARSAMNRHNNEAGROV
					1	GSPPARVLGKKEPPOOGVCPGS
						LGHMARSLAVGPQYSSLGSQPI
						LCASIPGLVPKQLRFCRNYVEI
			ŀ			MPSVAEGIKIGIQECQHQFRGR
			l	I		RWNCTTVHDSLAIFGPVLDKG
				I		YFGYLETITVPSDSVVSTKAETP
			I	I		DKGVCGRELGFRGGWLPELQS
1			l	I		LDEISMGWALCSRLOIKATRES
						AFVHAIASAGVAFAVTRSCAEG
		1	l	l		TAAICGCSSRHOGSPGKGWKW
		1		I		
			1	I		GGCSEDIEFGGMVSREFADARE
			l	L		NRPDARSAMNRHNNEAGRQ

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23375	53743	Α	23510	2094	2619	ASPASPAPCSGIPRT*SPYPSPSR
	l				TQ\APRARGPAIHAHPGP/RGCT	
			ŀ			ALPROGPPTPGSARAPPRIPAHR
						APTSPTLNFCLASSRQSMPPAPR
						PPPVPATNPVTPGCGRPPPTLGC
						RNPQTLRPPPGLPSPALLRRPLP
			ł			PPARVGMPPHQPPLPTTPACWR
						PVDWARAEGAGRAVC
23376	53744	Α	23511	3050	3317	PARTSPSASGPP**DLISALWTPS
						QPRAPTRQSLNSLSSSAPSSSTIG
						PSPSPPAPGPAPPLPWLPPSPSSL
						PRLSQVSRRPHSHLHPQ
23377	53745	A	23512	862	2954	AEHTLLVCFLNTSLTTTSTHVES
		l				FGGQLRGSQIGHPRRPNKMPYA
			l			WSIPQDPPLKPERDPGGPLGVS
		l				LSLPVNQYGESEEPMGDDDYD
					Į.	EYSKELNQYRRSKDSRGRGLSR
						GRGRGSRGRGKGMGRGRGRG
1						GSRGGMNKGGMNDDEDFYDE
1			İ	}	l	DMGDGGGGSYRSRDHDKPHQ
İ						QSDKKGKVICKYFVEGRCTWL
l						YHTTGNCINGDDCMFSHDPLTE
						ETRELLDKVEMTGSSVFDYIHP
			İ			GDHSEVLEQLGLRTPTPGPPTPP
					i	SVSSSSSSSSSLADTPEIEASLTK
						VPPSSLVQERSFFVRMKSTLTK
						RGLHVKASGYKVIHVTGRLRA
						HALGLVALGHTLPPAPLAELPL
						HGHMIVFRLSLGLTILACESRVS
						DHMDLGPSELVGRSCYQFVHG
						QDATRIRQSHVDCETHLHPPSL
						PTTPQTRASHSLVPGSPSSGLPT
						VPPAAPTAPGTPFSIGRAARRLS
						DASGLLDKGQVMTGYYRWLQ
			İ			RAGGFVWLQSVATVAGSGKSP
			1			GEHHVLWVSHVLSQAEGGQTP
		1	ĺ		I	LDAFQLPASVACEEASSPGPEPT
						EPEPPTEGKQAAPAENEAPQTQ
						GKRIKVEPGPRETKGSEDSGDE
						DPSSHPATPRPEFTSVIRAGVLK
						QDPVRPWGLAPPGDPPPTLLHA
						GFLPPVVRGLCTPGTIRYGPAEL
İ						GLVYPHLQRLGPG\PALPEAFYP
		l		l		PLGLPYPGPAGTRLPRKGD

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23378	53746	Α	23513	1319	2211	GSKVYGSLFHINHGNPFNMEVL
						VDSWPEYQMVIIRPQ/ETVRGE
						EKVQGNDLTFHHPTQEMTDDM
1		ŀ		1		DSYTNVYRVFSKDPQKSQEVL
	Ì	ŀ				KNSEIINWKQKLQIQGFQESLG
				l		EGIRAAAFSNSVKVEHSRALLF
		l				VTEDILKLYATNKSKLGSWAET
						GHPDDELESETPNFKYAQLNVS
l		l			İ	YSGLVNDNWKLGMNKRSLRYI
		1				KRCLGALPAACMLGPEGVPVS
ŀ		1				WVTMDPSCEIGMGYSVEKYRR
		l				RGNGTRLIMRCMKYLCQKNIPF
ĺ	1	l				YGSVLEENQGVIRKTSALGFLE
						ASCQWHQWNCYPQNLVPL
23379	53747	A	23514	234	858	SVPLQESPLPPPPTHQADC*SPC
		l		1		GAQGGWVTLAVASASS*AHTA
		1				SPLPRSPQAQGRAPSPAPPQIHL
						TTPSSFSSQGT/SHGGLGSTPGA
	ļ	1				DDPNTSSS\PPPGGSTPRWSAGD
	1	1				SAHSGGEVDGWS*SFP*SVPSY
ł	İ					GT*SPFCPWPSSGQRSFHGIASG
		1				WEPDGIPDGSSRWEPCGKPQES
		1				ETGVEGRQHFPKQLCLGSCRA
						VASPRVARA
23380	53748	Α	23515	1	3582	
23381	53749	Α	23516	628	4254	RLPESRAHDVAGADERPWPVH
						AGIFKSPAAGAAMSPAAAAAG
		1				AGERRRPIASVRDGRGRGCGGP
	İ	1				AGAALLGLSLVGLLLYLVPAA
	į.					AALAWLAVGTTAAWWGLSRE
		1				PRGSRPLSSFVQKARHRRTLFA
						SPPAKSTANGNLLEPRTLLEGP
1				•		DPAELLLMGSYLGKPGPPQPAP
		1				APEGQDLRNRPGRRPPARPAPR
		l				STPPSPPTHRVHHFYPSLPTPLL
					1	RPSGRPSPRDRGTLPDRFVITPR
		1				RRYPIHQTQYSCPGVL
23382	53750	A	23517	1	392	KVHMNGLRYALYIMGPALTRG
1		1			}	CHQVLIMLCCEVIERILT/VMM
1		l				KEAYFGEVDIMDDLPT\GNISA
1		1				NVTGRLYKCGVINHTVDLRRM
1		1			1	KT/EKWQNNLLPSRQFGFIVLTT
L	L	<u> </u>				SAGIMDHEEARRKHTGGKILGF
23383	53751	Α	23518	227	352	

SEQ ID NO:	of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino neid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23384	53752	A	23519	3	507	SFRAATMYRMNVLADALKSIN NAEKRGKRQVLIRPCSKVIVRF LTVMMKHGYIGEFEIIDDHRAG KIVVNLTGRLNKVGVQWRDLG YCNLCLPGSSDSSASASLVAEIT CGVISPRFDVQLKDLEKWQNN LLPSRQFGFIVLTTSAGIMDHEE ARRKHTGGKILGFFF
23385	53753	А	23520	279	653	ARAQRFSSQVDVFYLIQSVIFCR LALAEFEIIDDPQSLGKLL*TLT GRLKQ/CLGVISPRF\DVQLKRP WKKWHEIIL/PSNPAKFGF\IVLT TSAGLHG/HHEGRQDRKHTGG KILGILLYRDVIHIFT
23386	53754	А	23521	1	392	KVHMNGLRYALYIMGPALTRG CHQVLIMLCCEVIERILT/VMM KEAYFGEVDIMDDLPT/GNISA NVTGRLYKCGVINHTVDL/PPY ENGKWQNNLLPSRQFGFIVLTT SAGIMDHEEARRKHTGGKILGF
23387	53755	Α	23522	108	371	LSYIGKFEIIDDHRAGKIVVNLT GGLNKCGVISLRFDVQLKDLEK WQNNLLPSCQFGFIALTTSAGI MDHKEAR*KHTAGKILEFFF
23388	53756	Α	23523	252	378	
23389	53757	A	23524	3	518	HASAFRAATMVRMNVLADAL KSINNAEKRGKRQVLIRPCSKVI VRFLTVMMKHGYIGFEIIDDH RAGKIVVNLTGRLNKVGVQWR DLGYCNLCLPGSSDSSASASLV AEITCGVISPRFDVQLKDLEK WQNNLLPSRQFGFIVLTTSAGI MDHEEARRKHTGGKILGFFF
23390	53758	A	23525	279	709	ARAGRESSQVDVFVLIQSVIFCR LALAEF*NPLMTHQSLGKFVVN PHRAGLNKCGVISPQILTVQLQ RPWKK WAGLILLP\SRQVLGFI VLTTSAGI\MDHEGARPKTPRR GKSWGFFELDDVNYILFYKLKC LMDSGASTWSF
23391	53759	A	23526	207	564	SMAKEAISINRSLWAGPAKISSP AETEUESLISLFQKYYGKGGYN CTLSKTEFLSFMNAELAAFTKN QKDPGVLHRMMKKLGTNNDG QLDFSEFLNLIGGLAMACHDSF LKAVPSQKRT

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23392	53760	A	23527		407	FREKQSQELSQYGAMGAGQSF NSQFLQHGGPRGPSVPAGMNP TGIGGVMGPSGLSPLAMNPTRA AGMTPLYAGQRLPQHGYPGPP QAQPPAPTGGQENLL*GYSRA AVSARSLPGPFAIPVSHLAGNP TPPMNPH
23393	53761	Α	23528	3	315	
23394	53762	٨	23529		763	DYESTIOPTCSWKP/PVFKPDM IKIEEPOGPALKRCKTVSPAHV LMPSVMEMIAALGPGASPFAPL QPPSVPAPSDYPGQGSSFLGPGT EYGSPIPSSLTSEKSTSLPLQA RRPPAGHLDPTHNPGTPGLTN NLGAPPGPQLHHSNPPASRQS LGQASLGPTGELAFSPATOVMG PFSMSGAGEAPEPALDLLPELT NPDELLSYLGPPDLPTNNNDDL LSLFENN
23395	53763	A	23530		2901	MERRGPGAATARGRARPGGGP SVGLLATGSSLNPSFHGVARIVP GFIRIARPRDGSFAYESVPWQQS ATQPAGSLSVVTTVWGVGNAT QSQVLGNPMGPAGSPSGSSMM PGVAGGSSALTSPQCLQQAFA EGGANKGYVQQGVYSRGGYP GAPGFTTGYAGGPGGLGLPSH AARPSTDFTQAAAAAAVAAAA ATATATATATATATATALQEKQSQE LSQYGAMGAGQSFNSQFLQHG GPRGPSVPAGMNPTGIGGVM
23396	53764	Α	23531	80	412	KPGNGACAGREWCDGGGAAW NWRDPGLPVGDSGVWDRVLLE LLGPRSPRPLDVGGPAAGTPGV LSR/HLPVDCGLGPEALLCSSAP AARRPSLCGGHGRVMRGFRLG CGFRP
23397	53765	A	23532	1	192	RPLDVGGPAAGTPGVLSR/HLP VDCGLGPEALLCSSAPAARRPS LCGGHGRVMRGFRLGCGFRP

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23398	53766	A	23533	1	2094	RPLDVGGPAAGTPGVLSRPCPS TAALAPKPFCAAPRPQPDAPAC AGATGGSCADFDSGVDFVRR SSGLWGPQPPLFPVKNYTEMFQ DPVAFKDVAVNFTOEEWALLD ISQRHLYREWHLDTFWNLTSIG KKWKDQNIEYEYQNPRRNFRS VTEEKVNEIKEDSHCGETFTPV DDDRLNFGKKASPEVKSCDSF VCGEVGLGNSSFNMNIRGDIGH KAYEYQEYGFKPCKCQQPKKA FRYRPSFRTOERDHTGEFWAL LKNDGKKPLFYHSSIQRHMVV
						HSGDOPYKCKFCGKAFHWLSL VLHERTHTGEKPYECQCGKS FSYSATHRIHERTHIGEKPYECQ ECGKAFHSPRSCHRHERSHMGE KAYQCKEGKAFMCPRYVRRH ERTHSRKKLVECKQCGKALSSL TSFQTHIRUMHSGERPYECKTCG KGFYSAKSFQRHEKTHSGEKPY KCKQCGKAFTRSGSFRYHERTH TGEKPYECKQGKAHERTHTG EKPYECKLYGKALSRLISFRRH MRMHTGERPHKCGKAFSS
						PSSYGNEENSHIGERFYRUNGE GKAFTCFHPFQYHERTHTIGEK PDGCKQCGKAFRSAKYIRIHGR THTGEKPYECKQCGKAFHWVS SFHRHERTHAGEKPYECKHCG KAFTCSIYIRIHERIHTGEKPYQ CKECGKAFIRSSYCKKHERTHTI

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		L			1	
23399	53767	A	23534	48	1317	CGLMDELVHDLTSALEQTSEQ
	1	İ		1		NKLGELWEEMALSPRQQRRQL
				1		RKRRGRKRRSDFTHLAEHTCC
			l	1		YSEASESSLDEATKDCREVAPV
						TNFSDSDDTMVAKRHPALNAI
	1	1				VKSKQHSWHESDSFTENAPCRP
		1		l		LRRRR\KVKRVTSEVAASLOOK
						LKVSDWSYERGCRFKSAKKOR
	1					LSRWKENTPWTSSGHGLCESA
						ENRTFLSKTGRKERMECETDEO
	1					KOGSDENMSECETSSVCSSSDT
	1	İ		1		GLFTNDEGRQGDDEQSDWFYE
		l		ľ		GECVPGFTVPNLLPKWAPDHCS
				ľ		EVERMDSGLDKFSDSTFLLPSR
		1				PAQRGYHTRLNRLPGAAARCL
	l .	l				RKGRRRLVGKETSINTLGTERIS
		1		ļ.		HIISDPRQKDFWLPSAGKRERN
		l				QFNPLSPLYSLDVLADASHRRC
1						SPAHCSARQANVHWGPPCSRDI
		<u></u>				KQEAETSGHSIFV
23400	53768	Α	23535	225	403	
23401	53769	A	23536	1	403	REQEFLHMAPGEVTITVR\LIRS
		1				LEHR\NFKPVVYHGVNLDQTV
		1		ŀ		KEFIVFLKQDIPLRTNLPPPFRN
1		1		ļ.		YKYDALKIIHQAHKSKTNELVL
	1	1				SLEDDERLLLKEDSTLKAAGIA
						SETEIGIF\REEDYRNYKANPISS
23402	53770	A	23537	2	239	
23403	53771	A	23538	2	214	
23404	53772	Α	23539	1	1341	
23405	53773	A	23540	1	1677	
23406	53774	A	23541	1010	1509	
23407	53775	Α	23542	1	6355	
23408	53776	A	23543	223	10504	VKMPIGSKERPTFFEIFKTRCNK
	1					ADLGPISLNWFEELSSEAPPYNS
						EPAEESEHKNNNYEPNLFKTPQ
						RKPSYNQLASTPIIFKEQGLTLP
						LYQSPVKELDKFKLDLGRNVP
	1			1		NSRHKSLRTVKTKMDQADDVS
	1					CPLLNSCLS\ESPVVLQ\CTHVTP
					l	QRDKSVVCGSLFHTPKFVKGR
1	1	1			1	QTPKHISESLGAEVDPDMSWSS
1	1	1			1	SLATPPTLSSTVLIVRNEEASET
1	1	1	1		I	VFPHDTTANVKSYFSNHDESLK
		1			1	KNDRFIASVT
23409	53777	В	23544	137	709	
23410	53778	A	23545	1	1974	
	1-2.10	٠.	1-22.5	1'	I	

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23411	53779	IA	23546	207	1622	MACCL\SDEVKESKRINAEIEKA
23411	33777	1	25540	207	1022	AAAETRRDARREL/RRLLLLGT
	1	1	i		1	GESGKSTFIK\QMRIIHGAGYSE
		1				EDKRGFTKLVYQNIFTAMQAM
		1			l	IRAM/ETL/KILYKVRSRTRANA
		1				LILIRGGGTLEKVTTFGAFSYVO
		1				LPLKDLFGRDPGIQE/CAYERQP
	İ					RSYQLSDSAKYYLTDVDRIATL
		1				GYLPTOO\DVPAGSRVPTT\GIIE
		1				YPFDLENIIF\RMVDVGG\ORS\E
		1				RRKWIHCFENVTSIMFLVALSE
		1				YDOVLVESDNENRMEESKALF
						RTIITYPWF\QNSSVILFLNKKDL
						LEDKI\LYSHLVDYFPRSSMGPQ
						R\DAKAAREFILKMFVDLNPNS
		1				DKII/YTSHFTCPHRHRRTSRFVF
						PAVKDTILQLNLEEYKPGLSAP
						GPGRRDGDTGQDLPSTEPAAA
		1				RAGGAAESGPGPLPAGGDFFFF
		1			1	IFLTNGFYFTVIRGCTSLPPYTSR
		1				TFSPFVNGKGSLFLALTYGSLFS
		1				KKKKKKKE
23412	53780	Λ	23547	1	401	KKVPGRLSEMSFSLNFTLPANT
23412	33760	ľ^	23347	1'	1	TSSPVT\DCGPSLGLAAGIPLLV
		1		i		ATALLVALLFTLIHRRRSSIEAM
		1				EESDRPCEISEIDDNPKISENPRR
1		1				SPTHEKNTMGAOEAHIYVKTV
1		1				AGSEEPVHDRYRPTIEMERRR
23413	53781	A	23548	584	1071	PCRLPLHLILVEDLSFSLNFTLP
25415	33761	l^	23340	304	1071	ANTVSTAAPIHQAT/PEKTSAEF
i		1				TDCGPSLGLAAGIPLLVATALL
						VALLFTLIHRRRSSIEAMEESDR
						PCEISEIDDNPKISENPRRSPTHE
						KNTMGAQEAHIYVKTVAGSEE
						PVHDRYRPTIEMERRRGLWWL
		1			1	VPRLSLE
23414	53782	A	23549	25	654	LFLOMKDTSPIIFVRKELTIROC
	100	Γ.				TKKVPGRLSEMSFS\LNFTLPAN
		1				TTSSPVTGGKETDCGPSLGLAA
		1				GIPLLVATALLVALLFTLIHRRR
		1				SSIEAMEESDRPCEISEIDDNPKI
	1	1				SENPRRSPTHEKNTMGAQEAHI
		1				Y\VKTVT\KSEEPVHDRYRPTIE
		i				MERRRGLWWLVPRLSLGMMP
		ı				TSNQGSHQTWHLGTGLKTQGF
	1	1	1			VLGEERCOAAS
23415	53783	A	23550	108	413	RRSGSMALWRAYORALAAHP
1-55	1	1				WKVQVLTA/GLLGRKAALLAP
	1	1	l			CQ*LYLPTQASWTEGHASCSSL
			l	1		OEATSCPSAFCA*NSKATKOLV
		1	l	1		ERRGLQEHQRGRTLTMVS
	L		L	J	l	Diator delidiron in 12

SEQ ID	SEQ ID NO: of peptide	Met	SEQ ID NO: in USSN	Nucleotide location of first		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \-possible nucleotide insertion)
23416	53784	Α	23551	I	146	
23417	53785	Α	23552	2	442	GREARRSRSMALWRAYQRALA
	1					AHPWKVQVLTA\ASGEEGSTA
l						GPMPVAIPADPGLLD*RTRFLLL
						TPRSHFLPLCLLRMKLKSHKRS
l				l		LMGLGDIISQQLVERRGLQEHQ
l			İ			RGRTLTMVSLGCGFVGPVVGG
						WYKVLDRFIPGTTKVDA
23418	53786	A	23553	3	675	RHSRFGAQGSMALWRAYQ\RT
						LF\ALPWK\VQVLTAGSLMGLG
				l		DIISQQLVERRGLQEHQRGRTL
1						E/TMVSLGCGFVGPVVGGWYK
		i				VLDRFIPGTTKVDALKKMLLD
						QGGFAPSFLGCFLPLVGALNGL
1						SAQDNWGQTTAGELG/RDYPD
l			}			ALITNYYLWPAVQ\LANFYLVP
				l		LHYRV\AVVQCVAVIWNSYLS
l						WKAHRALSLPHSIVSTLQVMQ
					,	LDPWNGSDNPPQKWA
23419	53787	A	23554	512	1227	LQGHLMGIMAASRPLSRFWEW
						GKNIVCVGRNYADHVREMRSA
						/VWLSEPVLFLKPSTGVRAQRA
ļ						RPILMPAYTRNLHHELEL\ALVI
			1			GQ\RCRAVPE\AAAMDYVGG\Y
			l	l		ALCL\DMTAWDVQ\ASARKKG
						LAWDSGE/DAFTASCPVQRVSL
Į.						PKEKIPDPSQ\LKALGFKVNGEL
l			1			RQEGETSSMIF\SIPYIISYVSKIIT
			l			LEEGDIILTGTPKGVGPVKEND
						EIEAGHTRAGQYDHLKVEKAQ
23420	53788	Α	23555	192	436	RTSDHECFREGQICSVCLRERL
						NLNPEEACLLSQAPCMLFVLQY
				İ		LNTVIPYEKKASPPSV\EDLQML
l				l		\TNILFAMKEDNEKVP

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23421	53789	A	23556	94	1252	ARMEAPLVSI.DEEFEDLRPSCS EDPEEKPQCFYGSSPHHLEDPSL SELENFSSEIISFSSMEDLVNEF DEKLNVCFNVAKTENLAPV KNQLQIQEEETLQDEEVWDA LITDNYIPSLSEDWRDPNIEALN OKCSDTEIHEKEEEFNEKSEN DSGINEEPLLTADQVIEEIEEMM QNSPDPEEEEVLEEEDGGETSS QADSVLLQEMQALTQTFNNIW SYEGLRHMSGSELTELLDQVEG AIRDFWEELVQQLARRDEE EKEVKNSFFTVLUEVQNKQKIE
						QRELMKKRRKEKGLSLQSSRIE KGNQMP\LKRFSMEGISNILQSG IRQTFGSSGTDKQYLNTVIPYEK K\AYPPSVEDLQMLTNILFA\M KEDNEKVPYFA
23422	53790	A	23557	391	1169	GSFFFFKYFAYSPMPKNSKVV KRELDDDVTESVKDLLSNEDA ADDAFKTSELIVDGQEEKDTDV EGKSEVEDERPAWNSKLQYIL AQVGFSVGLGNVWRFPYLCQK NGGVVPFECEQSSATTYYWYR EALNISSSISESGGLNWKMTICL LAAWVMVCLAMIKGIQSSGKV NTYDRVTMSATVLALQEQWQT FSPIRSGNGHLTGSGAQRTPCRI RRGGSQLRVCDGGKQQWWTV SQSSARAITNTDQKSVQLQDLIE
23423	53791	A	23558	321	568	CLWCGASWPRRHSSPGRPAGD HRGDRP/EGPDTQGPGRPEPSTE PAFNPQSSLDQGGGPAAPAVD DDTRRGHGGVCGDGGAGEA
23424	53792	Α	23559	510	690	LRSPQ\RGVAAPPFLGSPVPFGG FRVQGGMAGQPSAPPALQTDD PYSKAHSVTPKSTPRVT
23425 23426	53793 53794	A	23560 23561	25 84	480 292	VRAVQSLHFHFPWRR/ISCPSES TRSIHVDRLPRRNSASLSLSAFP VSAGALINGEAWVRMGQVPIP KAP

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23427	53795	Α	23562	1138	1924	CRSSATWTPWTTTGCRGDLTK
						TYSLEAYDNWFNCLSMLVATE
		1		l		VCRLLLSPGSAMLRSPATFQVV
				1		KKKHRTRMLEFFIDVARECFNI
						GNFNSMMAIISGMNLSPVARLK
						KTWSKVKTAKFDVLEHHMDPS
						SNFCNYRTALQGATQRSQMAN
						SSREKIVIPVFNLFVKDIYFLPQ
	1	1			ŀ	NP\SNHLPNGHINFKKFWEISRQ
						IHEFMTWTQVECPFEKDKKIP\S
1		1		i		YLLTAPHPTARKLSSSPSFESEG PENHMEKDSWKTLRTTLLNRA
23428	53796	 -	23563	3	260	PENHMERDSWRILRIILLNRA
23428	53797	A	23564	1	1386	
23429	53798	A	23565	137	611	MKVNLGDSQNLACVFCRKHD
23430	33/76	1	23303	137	l ⁰¹¹	DCPNKYGEKKTKEKWNLTVH
						YYCLLMSSGIWQRGKEEEGVY
		1			1	GFLIEDIRKEVNRASKLKCCVC
	1					KKNGAFIGCVAPRCKRSY\HFP
				İ		CGLORECIFOFTGNFASFCWDH
	1					RPVQIITSNNYRESLPWHHLLG
	1				1	NLFEPIPSY
23431	53799	Α	23566	181	486	DPQGEAKGAAPAEGGGGPV*W
						N\GLQGPAGVPGRDGSPGANGI
	1					PGTPGIPGRDGFKGEKGECLRE
		1				SFEESWTPNYKQCSWSSLNYGI
						DLGKIAECTFTKMRS
23432	53800	Α	23567	600	888	QYNGMCSPGPAGVPGR\EGSPG
						ANGIPGTPGIPGRDGFKGEKGE
	1					CLRESFEESWTPNYKQCSWSSL
	1					NYGIDLGKIAGRNCWHILAPVP
		L				NIVSLYFGS
23433	53801	Α	23568	91	1170	ISLHRRPGPTCSMESPASSQPAS
	1					MPQSKGKSKRKKDLRISCMSKP
	1	1				PAPNPTPPRNLDSRTFITIGDRN
						FEVEADDLVTISELGRGAYGVV
	1					EKVRHAQSGTIMAVKRIRATV
						NSQEQK RLLMDLDINMRTVDC
	1					FYTVTFYGALFREGDVWICME
	1					LMDTSLDKFYRKVLDKNMTIP EDILGEIAVSIVRALEHLHSKLS
	1					VIHRDVKPSNVLINKEGHVKM
1		1	l			CDFGISGYLVDSVAKTIN\AGCK VKHGHESLN\PDLNRKGCDVKS
1			1			HVWSLGITMIEMAFLRFPYESW
						GTPFQQLKQVVEHPSPQLPAGR
		1		1		FSPEFVDFTAQCLRKNPAERMS
				1		YLELMEHPFFTLHKTKETDIA\A
1						FVKEILGEDS
23434	53802	В	23569	111	193	
23434	23802	ъ_	-5509	1	1.75	

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X-Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
23435	53803	A	23570	312	696	RP*SAHPEPGDCPSPPHPPRCGG VAQTAQVPDAPAPV*AG\QHSR SYMEDHLKNKNRLEKEWEALC AYQAEPNSSFVAQREENVPKN RSLAVLTSQPTRNQCLPFLQHG PGDICTPVSISSIVLCHRVT
23436	53804	A	23571		2974	QAAGMGPPLPLILLILLPPR VLPAAPSSVPRGRQLPGRLGCL LEEGLCGASEACVNDGVFGRC QKVPAMDFYRYEVSPVALQRL RVALQKLSGTGFTWQDDYTQY VMDQELADLPKTYLRRPEASSP ARPSKHSVGSERRYSREGGAAL ANALRRHLPFLEALSQAPASDV LARTHTAQDRPPAEGDDRFSES ILTYVAHTSALTYPPGBRTQLH EDLLPRTLGQLQPDELSPKVDS GVDRHHLMAALSAYAA
23437	53805	A	23572		3202	ASRRIPROHGLSAAGQAAGMG PPLPLLLLLLLLPPRVLPAAPSS VPRGRQLPORLGCLEEGLCGA SEACVNDGVFGRCQKVPAMDF YRYEVSPVALQRLRVALQKLS GTGFTWQDDYTQVMDQELA DLPKTYLRRPEASSPARPSKHS VGSERRYSREGGAALANALRR HLPFLEALSQAPASDVLARTHT AQDRPPAEGDDRFSESILTYVA HTSALTYPPGSRTQLREDLLPR TLGQLQPPELSPKVDSG

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	endon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23438	53806	Α	23573	I	1459	MPLTSQDVDQKTAAWVPTNRD
						STVPSDPWDCKRRPWPLLVLRP
	1		Î			LAPGTKAPPGGQNGRGRQKGS
	1		i			PCSRRLLLEPAGPLGPASVRCIR
						TTTPOLLIITTSLRFPSLEAAVGY
	1					GSNTCIFANACQMVFIMITWPY
1				i		LOCIPYSVLLKDLEMRNLRELE
					İ	
				1		DLIIEAVYTDIIQGKLDQRNQLL
1		-	ł	1		EVDFCIGRDIRKKDINNIVKTLH
				l		EWCDGCEA\ALLGIEQQVLRAN
1				l		QYKENHNR/TLQQQVEAEDVYI
1						LPQSGNDCRGLRFVSSLIGGVR
		1			ļ	CVHLKHKGDAVVWEAIEEGEA
					1	LPGKYRLKGGRSVWAAAVRA
						VCMDQVTNIKKTLKATASSSA
1			ŀ		ł	QEMEQOLAERECPPHAEQROP
ŀ			i		l	TKKMSKVKGLVSSSPLGPAGA
1			l	l.		AGTHQAWVRWGGDTKGPFPPL
				l		STCSEFOTCPSPHORLPTLLVLF
1					ŀ	
						QKNCYSPSPTPSFPSCSLQTQGL
						HQCHPKTGSDTAQLPSRRFLSL
						CKGLVSLPVFLLLPRHFVRLVIK
23439	53807	Α	23574	2	396	
23440	53808	Α	23575	1	1014	
23441	53809	Α	23576	110	212	
23442	53810	Α	23577	1	747	
23443	53811	Α	23578	323	495	GRAGKWHTQPE*SLRKAM*MF
1		1		1	İ	QILMMKMIVALARATSGTEAV
1		1	l	l		AGRSWASGLRRSGRV
23444	53812	A	23579	26	229	
23445	53813	Α	23580	I	574	
23446	53814	A	23581	889	1189	
23447	53815	A	23582	170	625	GAAVALVRPNGGTAASHVVCR
23447	1550.5	ľ.	20002	1	1020	EMRPLRPLOSRRCSRRRTGISN
		į.				MRALENDFFNSPPRKTVRFGGT
1	1	1	l	I		
1		1	l	1	1	VTEVLLKYKKGETNDFELLKN
1	1	1	l	1		QLLDPDIKDD\QIINW\LLEFRSS
		I		1		VMYLTK\DFE\QLISIILRLP\WL
		_				NRSQ\TVVEEYLAFLGNLVSA
23448	53816	Α	23583	166	455	RWKPSGTWPRNPMERPGH*ES
		l		l		VPKPGD*AKKSGGET\RGRKEG
1		1		I		NEHRETFSQIPDCHLNKKSQTG
		1	ĺ			VKPCKCSVCGKVFLRHSFLDRH
		1		1		MRAHAGHKRS
23449	53817	A	23584	3	3706	
	1-2011	<u> </u>	1	<u> </u>		l

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X-Unknown, *-Stop endun, /-possible nucleotide deletion, \-possible nucleotide insertion)
23450	53818	A	23585	11	1970	MVTDRKAAAGTHGEARPCWG
23430	33010	^	23363	1,	1970	SGSRVHTYRGAGERAMRVHRI
				i		MPREDSERVREADGGKHEORS
						FGDLHPTA APAPA ASGCGTQP
				İ		RVKQPGEWQHREAPQQRQPP\
		İ			1	RPHGSHQLVTVRYERRPGGKE
					1	RARLRPGLGCPSGRTPQGRAPG
	1				ŀ	GTALPTAAAAAIFPSAALTSGQ
	1	l				RNASLPKFYFRGGTWGTSQSE
		l				LCGKLSNQFLGEPLVPRIRLILS
		l				CGTCSGCSSPRKDPGHPEAGN
		l			1	PSRDFFQLPSEERDSVAFEDVA
		1				VNFTQEEWALLSPSQKNLYRD
		1				VTLETFRNLASVGIQWKDQDII
		l				NLYQNLGIKLRSLVERLCGRKI
	ŀ					GNEHRETFSQIPDCHLNKKSQT
	ŀ	ı				GVKPCKCSVCGKVFLRHSFLD
		İ				HMRAHAGHKRSECGGEWRET
	ŀ					RKQKQHGKASISPSSGARRTV
						PTSMRPYE\TKVCGKA\FIS\PIY
		ı				QIHQRTPTEKRSYKCREIVRAF
						VSSFFRKHGKMHTGEKRYECK
	l					YCGKPIDYPSLFQIHVRTHTGE
	İ					KPYKCKQFGKAFISAGYLRTH
	İ			l		IRSHALEKSHQCQECGKKLS\C
		l				SSLHRHERTHSGGKLYECQKC
		1		l		AKVFRCPTSLQAHERAHTGER
		1		1		YECNKCGKTFNYPSCFRRHKK
		1		j		HSGEKPYECTRCGKAFGWCSS
	ļ	Ļ				RRHEMTHTGEKPFDCKQWVK
23451	53819	Α	23586	86	390	LGSGDLPWEINPLSSCSLLREK
		l		l		PPTTSGPQT\TSPRNISPISNPEL
						THAGNLATGPRNARSPGFLLSI
		1		ŀ		VPSVWDPTENRTVQLTWQPLP
22152	50000	١.	22507	ļ	500	ELLELWPKVL
23452	53820	A	23587	1	502	MLLTQSLFGGLFTRTHMKFGA
		1				VTQIRGPPLGDKSPVLLLFALE
	1	1				QRRHVLSMDPKLRCWSRTGK
		1			1	AFPWCLIIAEMPDYSPTFQRCQ
		1				TTQGRLPWSFTLSSKSRFSGEG
		1				ARACYKCQKSDHQAKECPQPO
						IPPKPCPICAGP\TGNRTVQLTW
		⊢				QPLPEPLELWPKAL
23453	53821	В	23588	1	663	
	53822	Α	23589	233	496	
		IA.	23590	2	378	ERQRRHVLSMDPKLRCWSRTC
23454 23455	53823	1.	i .			
	55823	ľ				KAAFPWCLIIAEMPDYSPTFQR
	153823					CQTTQGRLPWSFTLSSKSRFSG
	53823					

SEQ ID	SEQ ID NO:	Mct	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23456	53824	Ā	23591	3	452	VRPRRDACLGPSPLAASPAFLG
				ľ		KGQVPQPLISLCPDPLFPHPNLIS
		l				LRPNPLCPHPDLVSLCPDPFPAF
						LEAHKNFOTPEPOOPGIPPEPPP
						PGACYKC/HEI*PPGOGMPAAO
1		l				DSS*AMS/LSVRDPTGNQTVQL
						TWOPLPEPLELWPKAL
23457	53825	A	23592	1352	1540	TWQTELEDESWITE
23458	53826	A	23593	177	383	EAMEPWKGVPLGRVNTEQIAP
23430	33020	r	23373	1'''	500	FSEAEGPREIH*GSRKVRKAAO
1						GLNEGPRELFWOPLPEPLELWP
		l				KAL
23459	53827	A	23594	1478	1744	ICAE .
23460	53828	A	23595	23	178	SCLEVCDEOGPEK\TRORALRG
23400	33020	ľ	23030	123	1.70	VSSVTEDTLNICRLCWOPLPEPL
		ļ				ELWPKAL
23461	53829	A	23596	1	282	EEWIKAE
23462	53830	A	23597	34	241	SGDLPWEINPPSSCSLLREKDPP
23402	33030	ľ`	1337,	1	1	MTSGPQT\TSPRNISPISNPDPTG
		l				NRTVOLTWOPLPEPLELLPKAL
23463	53831	A	23598	121	286	AASFYSSPTSLTIPOPLSPFNLGV
23403	33031	ľ``	23370	1.2.	200	TLOSLPSLNFNSF\LFWOPLPEPL
		1				ELWPKAL
23464	53832	A	23599	144	488	GYEIWCRDSDOGTFPWEINPPS
23404	33032	ľ.	23377	1	1.00	SCSLLREKDPPTTSGPOT\TRPR
				1		NISPISNLVSGLFLLSSPPSLTIPO
		l				PLSPFNLGATLQSLPSLNFNSFH
		1		1		FLVETKETRFIRGAKIPAPVTD
23465	53833	A	23600	471	677	TEVERNETIA INGILIATI II TIE
23466	53834	В	23601	67	555	
23467	53835	Ā	23602	1	1440	
23468	53836	Ā	23603	860	1209	IPGNCRDSSGVGIKERETNAGS
	1	1.		1		OHMH/IKNSSVHGYPRLCVEDE
		l			1	DAYKKOFSOYIKNSVTPDMME
		l		1		EMYKKAHAAVPENPVYEKKPK
	1	l				KEVKKKRWNRSKMSLAOKKD
1		1		1		WVAQKKASFLRA
	<u> </u>	L	L	1	L	THE PROPERTY OF THE PARTY OF TH

SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23469	53837	A	23604	1	966	MGFVKVVKNKAYFKRYQVKF
1	ł	1				RRRREGKTDYYARKRLVIODK
1		1				NKYNTPKYRMIVRVTNRD\IICC
1						IANA\RIEG\DMIVCAP\YAHELP
		1				KIWV*RVGLTNYAAA\YCTGLL
		1				AGPAGFFHRFGMDQIYE\GQV\E
		1				LTGDEYNVESIDGOPGCLHPAY
		l	1			WDARPLPRTYPLAIKVFGCP*R
		l				GA\VDGGLVLFP\HSTKTIPPGY
		l				DS\ESK\EFNAES/IHRKPHPWAQ
		l				NVARLHAPTLMEEDEDA/YTK
		l				KQFSQYIKNSVTPDMMEEMV/Y
		l				RKAHAAIRENP\VYEKKPKKEV\
		l				KRKRWNRPKMSLAQKKDRVA
		l				PKKASFLQSSRSGLLESLTPAIFF
23470	53838	A	23605	1	644	ANSSPVNPVVFFDVSIGGOEVG
		Γ.		ľ		RMKIELFADVVPKTAENFRQFC
		l				TGEF/RVSFRCV*LLCLLORGVL
		1				GLO*GYR*TTCWP*MIAGDSDR
				1		RKDGVPIGYKG\STFH\RVIKDF
		l				MIQGGDFVNGDGTGVASIYRG
		l			i	PFADENFKLRHSAPGLLSMANS
		l				GPSTNGCQFFITCSKCDWLDGK
		1				HVVFGKIIDGLLVMRKIENVPT
		l				GPNNKPKLPVVISQCGEM
23471	53839	A	23606	638	1391	CPAYKGAHGIQCVYCTLPPPLL
						QTTRRVRRNPTLPTSASGSEPW
		l				RLANSSPVNPVVFFDVSIGGOE
						VGRMKIELFADVVPNTAENFR
1		l				OFCTGEFRKDGVPIGYKGSTFH
		l				RVIKDFMIQGGDFVNGDGTGV
		l		ł		ASIYRGPFADENFKLRHSAPGL
1		l				LSMLTWPGKLTAGIFGFHQANS
		l		İ		GPSTNGCOFFITCSKCDWLDGK
		l				HVVFGKIIDGLLVMRKIE\GMPC
1						IFQPTWTHFLPVITTACSALGSM
1						LAAERGTKN
23472	53840	A	23607	2	591	ARAEFGTSRVGAMAVANSSPV
		l				NP\VVFFDVSIGGQEVGRMKIEL
		ı		1		FADVVPKTAENFROFCTGEFRK
		l		l		DGVPIGYKGSTF\HRVIKGF\MI\
		1		I	Ì	QGGDFV\NGDGTGSRPVFTRGP
		l		l		ILQMKIFKL\RHSAP\GLLSMAN
		1				SGP\STNGC\QFFITC\SKCDW/V
		1		l		WDGKHVVFGK\IIDGLLVMRKI
	l	1				E\NVPTGPNNKP\KLPVVISQCG
23473	53841	A	23608	6	424	10000
	1000.1	1		I*		L

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide togation of last	Amino acid sequence (X=Unknown,		
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide		
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)		
				sequence				
23474	53842	A	23609	3	470	DAWVSGRLTELPSSCFFPPIFRO		
				Ī		LLPVMPEPSKSAPAPKKGSKKA		
						VTKAOKKDGKKRKRSRKESYS		
i						VYVYKVLKQVH\PDTGISSKAM		
						GIMNSF\VNDIFE\RIAGEASRLA		
l						HYNKRSTITSREIQTAVRLLLPG		
1						\ELCKHAV\SEGTNA\VTKYTSS		
23475	53843	A	23610	93	42 I			
23476	53844	A	2361I	2	1173	DHAAPRRSSRASVATVTRAGR		
						CPTMSVRVARVAWVRGLGASY		
l		1				RRGASSFPVPPPGAQGVAELLR		
l		l				DATGAEEEAPWAATERRMPGQ		
		l				CSVLLFPGQGSQVVGMGRGLL		
ļ					1	NYPRVRELYAAARRVLGYDLL		
						ELSLHGPQETLDRTSQVQSPGT		
		1				GIPRRVRAHVLHLSPQVIENC/V		
						VS/AGLAFSPSFCALIFPNALL/F		
						WPGLYAVKIRAEAMQEASEAV		
						PSGMLSVLGQPQSKFNFACLEA		
Į.						REHCKSLGPVNGQICKGGFLGH		
						CSAETSFLPQALRFLQKNSSKF		
						HFRRTRMLPVSGAFHTRLMEP		
						AVEPLTQALKAVDIKKPLVSVY		
					İ	SNVHAHRYRHPGHIHKLLAQQ		
					İ	LVSPVKWEQTMHAIYERKKGR		
		<u> </u>		105	105	GFPQTFEVGPGRQLGAILKSCN		
23477	53845	A	23612	187	425	Maria de la porte de la composición dela composición de la composición dela composición de la composición de la composición de la composición dela composición dela composición de la	1258	MKAYLLLGAAPPLPWFPGQAV
		l				FRPCPPPAGLWRTAQLVELEHG PVPREDLIPCSRPFSISFLGPERV		
		l				DGLAFPRIQALGIAAARGMEPK GAPGKSDSGNSRSAGGRVLGC		
		l				OGAALEVECSAGRGGPVRLAV		
		i				GCVLLRAPERGWRRGEGAARG		
						IPAGRSSESLGALSPQVGVRSDI		
						GINGNIWHPGVOTDFSSLKRSO		
						GLYAVKIRAEAMQEASEAVPS		
						GMLSVLGQPQSKFNFACLEARE		
						HCKSLGIENPVCEVSNYLFPDC		
			1			RVISGHQEALRFLQKNSSKFHF		
						RRTRMLPVSGAFHTRLMEPAV\		
		1	l	1		EPLTQALKAVDIKKPLVSVYSN		
			i			VHAHRIQGIPGHIHKLL\AQQLV		
			l	1		SPVKWEQTMHAIYERKKGRGF		
			ĺ			POTFEVGPGRQLGAILKSCNMQ		
					1	AWKSYSAVDVLQTLEHVDLDP		
	L				l	IN WAS I SAVDVLQ I CERVOLUP		

SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nuclcotide
sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
			sequence		
53847	Α	23614	2	1050	GRCPTMSVRVARVAWVRGLG
					ASYRRGASSFPVPPPGAQGVAE
					LLRDATGAEEEAPWAATERRM
i					PGQCSVLLFPGQGSQVVGMGR
					GLLNYPRVRELYAAARRVLGY
					DLLELSLHGPQETLDRTVHCQP
					AIFVASLAAVEKLHHLOPSVIE
					NCVAAAGFSVGEFAALVFAGA
					MEFAEG\LRFLQKNSSKFHFRR
					T\RMLPVSGAFHTRLMEPAVEP
			1		LERKLLKAVDIKKPLVSVYSOR
					PTRIRLOAFPGNIOOSCLAOOLG
		-	1	Į.	LPK*KWEQ\TMHAIYERKKG/R
					RGFPQTFEVGP\GRQLG\AILKS
					CNMOAWKSYS\AVDVLOTLEH
					V\DLDPQEPPKMTGRGLKCNDP
ŀ					LCPPEERL
53949	Δ	23615	600	905	OAPASGSPWLSGGPOOVEDAG
33040	ľ	23013	000	J~~	AGYGFAPGOPPPPRTOPRSAC
					SRRAAG/TPVPRAPPAPSAG\PK
l	1				ARAPRSLGCGSLGRFSTGVRPD
ł					KCIFPPETENAAGCF
53849	A	23616	390	722	QAPASGSPWLSGGPQQVEDAG
			-		AGYGFAPGQPPPPPRTQPRSAC
					SRRAAG/TPVPRAPPAPSAG\PR
					AAAPRSLGCGSLGRFSTGVRPD
					KCIFPPETEMSYLLFRNTVSRCR
					c
53850	Α	23617	1	673	MGQVKGHKLCLDFTLLSPDGG
					AAEGRCLELGWOAKEVEETIE
1					GMLLRLEEFCSLADLIRSDTSQI
			i		LEENIPVLKA\KLTK\MRGIYAK
					VDRIKAFVKMVGHHVAFLEAN
					VLQAERDHGAFPQALRRWLGS
					AGLPSFRNDPTAPLRQVFEASP
1					DPRLARTVTIQTQQPVRVSGPL
					SGQRKEKSLQKGAQVFQTNVP
					ETLLEGAAISPAPACGGPLTSGD
					GFRHVRK
53851	Α	23618	1	651	
53852	Α	23619	195	497	DQFLENTKKCDGLNQLSFDPA
	l				NNNKPLEFGRARGPLVTEGLIE
	l				NGGESSKKRRRTNSSSADNSRT
					LNVDSTAMTLPMSDPTAWATA
1	1		1		M\NNFGMAPVRIVGH
	53847 53848 53850	53847 A 53849 A 53850 A	of peptide sequence bod in USSN 809/540,217 53847 A 23614 53848 A 23615 53849 A 23616 53850 A 23617 53851 A 23618	of peptide sequence bod in ISSN 109/840,217 location of first sequence 53847 A 23614 2 53848 A 23615 600 53849 A 23616 390 53850 A 23617 1 53851 A 23618 1	In USSN location of first codon for last amino acid sequence

SEQ ID		Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last aminu acid	*=Stop codon,/=possible nucleotide
	sequence	l	09/540,217	eodon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23485	53853	Α	23620	241	1823	NDDRAKFTVEDHTMQRDFRW
						LWVYEIGYAADNSRTLNVDST
						AMTLPMSDPTAWATAMNNLG
		ĺ				MAPLGIAGQPILPDFDPALGMM
						TGIPPITPMMPGLGIVPPPIPPDM
		1				PVVKEIIHCKSCTLFPPNPNLPPP
		l				ATRERPPGCKTVFVGGLPENGT
		ĺ				EQIIVEVFEQCGEHAIRKSKKNF
						CHIRFAEEYMVDKALYLSGNR
		l				ELKKFLNKEDNYFRKINLTALY
		l				KTVLEERCEKRRGGYRIRLGSS
						TDKKDTGRLHVDFAQARDDLY
		l			į.	EWECKQRMLAREERHRRRMEE
						ERLRPPSPPPVVHYSDHECSIVA
		l				EKLKDDSKFSEAVQTLLTWIER
					l	GEVNRRSANNFYSMIQSANSH
		l				VRRLVNEKAAHEKDMEEAKEK
						FKQALSGILIQFEQIVAVYHSAS
		1				KQKAWDHFTKAQRKNISVWC
		l				KQAEEIRNIHNDE\LMGIRREEE
		l				MEMSDDEIEEMTETKETEESEL
		l				KKLEGIVDGSVEVSAEALRKLL
		i				NEYERQVKEEQKVGYRIWGTI
						HNKGEGKKEEFAEELEKSSSER
23486	53854	Α	2362 I	996	I512	RPVHLASPAKEQQGTRKGRRA
		l				PARLPVPTTGRGLEKGPESQAP
		i			l	PSFRQATGRFSRSGSSGGPTPSK
		l				NSRDCSHTPQLPTPRGNPYVFPS
			ĺ			PGHPEPSRPVEMPAPG\EPRTRQ
		1			i	LCRERTRKACPPKPRPPLGLPG
						DPTGPVTHHAPPVSPTGASGQE
			l		1	RRAEPGAVSYAHASATK
23487	53855	Α	23622	293	1140	PRGILHQDKNLVVINK\PYGLPV
						HGWPWG\QLCITDVLPILAKML
		1				HGHKAEPLHLCHRLDKETTGV
		1		ŀ		MVLAWDKDMAHQVQELFRTR
		1				QVVKKYWAITVHVPMPSAGVV
		l				DIPIVEKEAQGQQQHHKMTLSP
		1				SYRMDDGKMVKVRRSRNAQV
		l				AVTQYQVLSSTLSSALVELQPIT
						GIKHQLRVHLSFGLDCPILGDH
		l				KYSDWNRLAPQKLSVGTLKKL
		l				GLEQSKARYIPLHLHARQLILPA
		1				LGSGKEELNLVCKLPRFFVHSL
		1	1			HRLRLEMPNEDQNENNEAKCL

SEO ID	ISEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X-Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
		1	i	sequence		
23488	53856	A	23623	1	1328	MAVDLGSHTPGSSGVTAATFR
23488	22820	<u>۱</u> ^	23023	1'	1328	
1	l	l				GRRLVNTREESVVLGAGGEVL
		1				SGVPGTWDALGPGASFGIKGM
		1				FPKGTVIMSTSPMCEGPVPRRK
		1				PCASSVGQQNSRSRKKAGRKTP
		1				TLAGRHVPVKIKKEAIQQKPLP
						ASSFSPALTRSGAIFGPCPALTS
		1		ļ		APVLEYPLQKVVENEAFKVFN
		1				NRDEESKRQKQAEFRMLTSAIR
			1			GPAGPRGHSSTRKPPSNPPPPGA
		l				CFKCGNEGHRSRQCPNPGTTSQ
	1	1	i			EAELIALTRALTLAAGQQINIYS
		l				NSHYVFHIVHSHSSIWKERGILT
	1	l				AKNTPAINGSLINKLLKAARLP
		l				QKVAIIHCRGHQTPDNPILAGN
		l				VLADQVAKQVALQPVQGQFLS
İ		1		1		LSLFSPLYSSEEKEDFRAQNLQ
ĺ				1		KQGPW/LQSQGARLLVPAACAS
	1					KVAEAGRKPAGRHVPLKIQKE
	1	1				AIWSSPFSVPMATRAAASTSPT
23489	53857	A	23624	475	622	IHSKVLFSASSSSC*NTNNSWFR
		1				ASIKKNPEKONKNMOSAIPOHG
		1				LWKI
23490	53858	Α	23625	17	992	GRGEVLRGRGRGQLDAVLSRE
1	1					HCGGSRQSEQPWSRRSSSSSSR
		1				RRLRSAAEPAMALSMPLNGLK
		l				EEDKEPLIELFVKAGSDGESIGN
i						CPFSOG\LFMILWLKGV\VFSVT
İ			i			TVALKRKPADCQNLVPGTPPP\
						LILFSSEVKTDVNKIEEFSEEVL
		İ				CPPKYLKLSPK\HPRSNTAGMDI
						FA\KFSAY\IKNSRPEANEALER
1	1	l				GLLKTLQKLDEY\LNS\PLP\DEI
1	ŀ	l				DENSMEDIKFSTRKFLDGNEMT
1	1	1		1		LADCNLLPKLHIVKVVAKKYR
1					1	NFDIPKEMTGIWRYLTNAYSRD
1	1	1				EFTNTCPSDKEVEI\AYSDVAKR
		ı				LHOVKSRLLKEVSFMSSP
1	I	ı	I.	I	1	LHQVKSKLLKEVSFMSSP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
23491	53859	A	23626	11	1128	MALASGEGLPVALSHGRRGNG
		i .		ľ		OERVPYLKLRDTCAYRVAWRC
						NRLARAKVKRTSYRSGYGNGF
		1	l			EPGVEPDCVVLLALGVENDLV
				l		RISSKAEKYSLYEFSFPLRVSRE
İ		1	l			CEEECRKEPWFNPGOKPSVLIT
		l	1			SGQILGQEEQKTLLFPGLLSLVF
1		1				OFNVENSDLRCLSLAENLEISAL
1		1				LSLSCSSVIRLILINLEVTISPPITI
1		1	1			
1			1			LKHAPPLTSLEASSSSSGIEKKIH NNFROGPFPSSATRSTFLMNHT
1		l				SFDSYVVVSNIVPRLNLYPOVL
		i				
		1				DVLQSLHKIHQLASNAKWTIQG
		l	i			PAKPDLGLELGLHRYPDWEKE
1		1				ERKKKRKRSVSPEKSITTHNKP
1		1				TSSGIFY/RPRRCGCDLGRPPAP
		Ļ.				GPHPSSPPHSPPQPSRCPPV
23492	53860	Α	23627	3	453	SLGKDIRPRSARAACKGVGLWS
		1		l		GCFGKMAGSGVRQVTSTASTF
		ĺ		1		VKPIFSRDMNEAKRRVRELYRA
		1				WYREVPNTGER*RLVRPAVLK
		_				HSYVANRSF
23493	53861	Α	23628	3	695	PYASRRKFRVDPRVRAWSGCF
		ı		1		DKMAGSGVRQATS\TASTFVKA
		l		1		HFSVGDMNE\AKREGARVLTAP
1		ł		1		WYR\EVPNT\VRVHQFQLDI\TV
		1		1		KMGRDKVREMFMKNAHV\TDP
1		1				RVVDL\LVIKGK\IE\LEETIKVL
		1				KQ\RTHVMRFFH\ETEAPRPKDF
		l				LSKF\YV\GHDPLKSFSGKDAR
		ł				WILFLGAQIKLTIQWSLCDRIPL
l		l		1		MNQFSAASFCLSKWGPWVLHI
		L_				TCSFPFSWKCGVCCKLD
23494	53862	Α	23629	82	441	SLPINPVPTACTLQPAGLLRPPG
				1		SDVRTPMLPPSAPSQPSYPTSRP
1		1				PLWPTWPAGPVFRRSQPLAQPN
I		l				LPRSYHQQAPGGQTPVALGLE\
		ł				CPKATNQPFGTLQRDPKSPAKV
		1				PPFEEFSL
23495	53863	Α	23630	3	504	DRSSSSTMSTGGDFGNPLRKFK
		ł				LVFLGEQSVGKTSLITRFMYDS
	i	1				FDNTYQATIGIDFLSKTMYLED
1		l				R/TNVNSFQQTTKWIDDVRTER
		1				GSDVIIMLVGNKTDLADKRQVS
1		1				IEEGERKAKELNVMFIETS/AKA
				1		GYNVKQLFRRVAAALPGIMPS
		1		1		VLFQIIPSQALTRYLC
	<u> </u>				L	. 5. 4 54

SEQ ID	SEQ ID NO:	Mct	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino seid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	delction, \=possible nucleotide insertion)
				sequence		
23496	53864	A	23631	416	1239	GGGRSSSSTMSTGGDFGNPLRK
						FKLVFLGEQSS\GKTSLITRFMY
	l					DSFDNTYQATIGIDFLSKTMYL
		1				EDRTIRLQLWDTAGQERFRSLIP
						SYI\RDSAAAVVVYDITNVNSFO
						ANYKSGIDDVRTERG\SDVI\IM
						LVGNKTDLADKRQVSIEEGERK
	1					AK\ELNV\ILMETSAKAG\YNVK
						QLLSTCKQPALPGMESTQDRSR
		ĺ				EDMIDIKLEKPQE\QPVSEGGLF
						LTNLPCHLQPFFQKLTGFWPPY
1	ŀ					SFIDCSVNIGLMPFPFSNNVLRFI
		1				IAACPAWR
23497	53865	A	23632	1	493	PKPATAPPPPSL/RARFLCPORRP
23497	33863	<u>۱</u> ^	23032	['	493	ERAR/PRARGASPRPEKTAGAR
i						KRRGSCERPHPGG/RPPAPG*I
	1					SSRPTLGFFWILNFFFLKKLGRI
						KDVPWODSTKEHLSVCRGLRG
1						TWSVPSASDPLHPAWRRTCFRV
1				1		PPTTGAREWWLHLPPPALORPP
		1		1		DSROCHRIPVC
23498	53866	A	23633	1	560	MESIWAGLWTAODTTVLADDF
23498	22800	I ^A	23033	'	300	EFILWTLKRRPERAR/PRARGAS
		l	1			PRPEKTAGARKRRRGSCERPHP
		1				GGLRRLPGARDQRAPPRSDAG
		l				RSPGRLRRLPALLCAGPLSPGRE
		l				PRRWEPRAGVARPAGQQWGFF
						WPGNQQGASICSLILWFLLECG
		1				SRSTGGPCSGTELGHFSRERPTV
		1	·			LLSRLGTVKRP
23499	53867	A	23634	32	327	LTORLFGGLFTRTHMKFGALTR
23499	33807	 ^	23034	32	321	IG\DLPWE\INPLSS\CSLLPEKDP
1			i			PMTSGPQTDQPKEHLTNFKSAA
		l				RRSSRTSSHRNLLHVLEIWOLD
						QGMPAARDF
23500	53868	Ā	23635	2	424	LTPFSDSAHLHPGEINSPVAHTK
23300	33800	^	23033	<u> </u>	424	PVGGLFTRMRVTFGAKTODRS
1		l	1			TP\QETSPLSSPS\PVTSGPQTNQ
1		i				PKEHLTNFKLGKQSFHSSHASL
1		l		l		ATLOSSSLATLOSPCPSNSNSFS
1						SLVETRRRPGAVARLQSQLSGR
						LRGGRIA
22501	62000	ļ	22626	31	246	
23501	53869	A	23636	Pi	240	TALLLTQSLFGGLFTRTHMKFG AVTRIG\DLPWGINPLSSCSLLR
	1		1	I		
	1		1	I		EKDPLTISGPQTHQPKEHLTNF
L		L	L	L	L	KSGPH

	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X=Unknown,
	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	coden for peptide	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
23502	53870	Α	23637	125	464	TAVFLTQSLFGGLFTRTRMKFG
						TMTRI/RGDLPWEINPLSSCSLL
						HEKDPPTTSGPQTNQPKKHLTN
						FKSRKVERTKGLLKTHLTKLSH
						QLKKAWTILLPLSLLRIQACPR
						NAT
23503	5387I	Α	23638	1064	1304	TAMLLTQSLFGGLFTQMCMKF
						GAVTRIG\DLPWEINPLSSCSLL
						HEKDPPTTSGPQTDRPKEHLINF
						KSGACYTCRKSGH
23504	53872	Α	23639	539	747	TAMLLTQSLFGGLLTRTRMKF
		ł				GALTRIG\DLPWEINPLSSCSLL
				1		HEKDPHTPSRKIHIRPQGNHESQ
						KGL
23505	53873	Α	23640	709	950	TALLLTQSLFGGLFTRTHMKFG
						AVTQIG\DLPWENQSPVLLCSLL
1 1						REKDPPTTSGPQTDQPKEHLTN
						FKSGKWSFHSSPAF
23506	53874	Α	23641	1	1257	MGKKVLGIASQLLGAVRCEPD
		l				GRRVLPEDWPECLSLVTHGTW
						FTRQGDKNIQMADNSFSDGVPS
						DFVEAAKNASKTEKLTDQVMQ
						NPGVLAALQERLDNAPHTPSSY
						IETLPKAVKRRINALKQIQVRC
						AHIEAKFYEEVHDLERKYAAL
					ļ.	YQPLFDERREFITGDVEPTSDM
			1	l	1	ESEWHRENEEEEKLAGDRKNK
						VVITEKAAATAEEPNPKGIPEF
						WFTIFRNVDMLSELVQEYDEPI
		l				LKYLQDIKFHFQPNDCFANSVL
		1		ŀ		TKTCKMKSEPDKADPFSFAGPE
		1				IMDCDGCTIDWKKGKNVTVKT
				l		IKKKQKRKGPGTVRTITKQVSN
						ESFFNFFNPLKASGDGESLDKD
		İ				SEF/ASDFEIGHFFRER\PQAVLY
						FAGEALKDDDNFDEGEEEEDEL
						EGDEEGEDEDEVEINPNKEPSQ
	53875	Α	23642	132	1277	
23508	53876	Α	23643	1	515	RVRHRSVSVGSTHASAHAYGE
						GSLRIIPVTRSPLVTMNHIVQTF
						SPVNSGQP\PNYEMLKEEQEVA
				l		MLGAPHNPAPPTST\VIH\IRSET
				ĺ		SV\PDHVV\WSLFNT\LFMNTCC
				l		LGFIAFAYS\VKSRDRKMVGDV
						TGAQAYASTAKCLNIWALILGI
				L		FMTI\LLIIIPVLVVQAQR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	coden for last amine acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23509	53877	A	23644	243	643	NHTVQT\FFSPVNSGQPPQRMR
						CIKEEHEVAVAGG\PHNPAPPTS
	l	1			ŀ	T\VIHIRSETSVP\DHVVWSRVQ
						TPSSMNPC\C\LGFIAIRLTP*SLR
						DRKMVGDVT\GAQA\YAST\AK
						CLN\IWAHDSGASVMTILLIVIPS
23510	53878	A	23645	3	1498	RCNFLRSSRIRVHPTPAASTMPP
					l	KFDPNEIKVVYL\RCTGGEVGA
	İ	1				TSALAPKIGPLGLSPKKVGDDIA
	l	1				KAMGDWKGLRITVKLTIQNRQ
						AQIEVVPSASAMIIKALKEPPRD
	l				ľ	RKKQKNIKHSGNITFDENGNIC
	İ					STDMR/HRSLSRELSGTTKEILG
						TAQSVGCNVDGRHPHDIIDDIN
	İ					SENDKQPRKSKMRTSFSPQIPG
						HEHFHYLLLLQHVHCQLALTL
						RARKEGSSTHHQKLSTYYFKEG
	l					NSKEPKKIQRCRNTELRKEEGG
	l	ļ			l	GGSGKAPGSEIKLESPACSGASL
						GEMTAWTMGARGLDKRGSFF
		1				KGPLPNTCGHFWEMVWEQKSR
		1			İ	GVVMLNRVMEKGSVVEFVTEL
	l	1				FFYVALPKMFPVHYIVILLASKL
						EEVTFNFPPPPDLCTPFLQNYLR
	i	i				VAFQEVNSGCTGKTLLVRPYIT
	i					TEDVCQICVEKFKVGDPEEYSL
					ŀ	FLFVDETWQQLAEDTYPQKIKA
		1				ELHSRPQPHIFHFVYKRIKNDPY
						GIIFQNGEEDLTTS
23511	53879	Α	23646	361	918	GCTGSRGTVPLSALGPQRIGPP
						GSCLQIKSWVNDICQRQPG\DW
			i			KGP*GFTVKLTHSRTGQAPDCT
		1				RVP/VLPSCP*FIQSPSREPPKRPE
		1				KKPEKTFKHQLGIFTFLNEICST
		1	i			LLRQMR\HRSLSPEKLSWNPT*
		1	l		i	RDSWGIAQFSGAVNVDG\RH\P
		I				HDI\IDDIQQWVLWKCPRQLST
		1		_		KENIFNKGII
23512	53880	Α	23647	1	642	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23513	53881	A	23648	894	2438	VETKRIQPQPPDEDGDHSDKED
23313	33001	l'`	250.0	0,,		EOPOVVVLKKGDLSVEEVMKI
İ		ı				KAEIKAAKADEEPTPADGRIIY\
		1				RKPV\KHPSDEKYSGLTASSKK
		1				KKPNEDEVYQDSVKKNSQKLII
İ		1				NSSLLSFONEDENDTEKSMENL
		1				TGKIRKLQLRIVDPTFLPKLSNK
		l				GKDPVAVARLETGKGNKGAFR
ŀ						VLRDISRWATLCTSPQHSSSEA
						KNIIPYQATQFQNEVHKWYKK
		1	l			KGGLKISKRKALFFLTQDIQKK
		l				LOKLALGPNTPTPDILKIASLVF
1		l	l			YNODQEEKERAQQNERQKEKL
1		l				YWLSYKSASPFQVALRILSQET
		l				THKDELPTDLPNNPEVQAPGIP
		l				GNTIMAMPVAIQLKDPSSYPCR
1		1				IIKVPEIASWIHRTQVKLWKSPE
		1	ł			TSELEPTPSVIEYTREALEDLKF
ł		1				LFKOKDNSSAKTPGRAYAASPR
						DWTSIEAELHIFYOWDPNLKG
		1				MFKPASLLAKVKODFPDIHKES
		l				PIFRAIFSNITLMGIAPICVTGKTI
i				}		
1	ĺ	i				KKRTRQHHTTPKSNLREQASQ GSLNWEGN
23514	53882	A	23649	200	1727	RTATSLDSVLYKSDYMTWKYR
23314	33002	ľ.	23047	200	1''-'	VSFFYILSCRLLKK/QCTDNEHV
						YMELSKKLYKYCPKEWKKEAS
	1				İ	KVRQYEVTWVVSKRGKDYILK
						HIPNMHKDOFALTASEAHLKYI
						KEAVRLDDVAVHYYRLYKDK
l			i			REIEASLTLGLTMRGIQIFONLD
		ŀ				EEKQLLYDFPWTNVGKLVFVG
l						KKFEILPDGLPSARKLIYYTGCP
		l				MRSRHLLQLLSNSHRLYMNLQ
		l				PVLRHIRKLEENEEKKOYRESYI
		1				SDNLDLDMDQLEKRSRASGSS
	l					AGSMKHKRLSRHSTASHSSSHT
						SGIEADTKPRDTGPEDSYSSSAI
						HRKLKTCSSMTSHGSSHTSGVE
I		1	l			SGGKDRLEEDLQDDEIEMLVD
1		1	l			DPRDLEOMNEESLEVSPDMCIY
I		1	1			ITEDMLMSRKLNGHSGLIVKEI
		1				GSSTSSSSETVVKLRGOSTDSLP
1		1	l			OTICRKPKTSTDRHSLSLDDIRL
I		1	l			YOKDFLRIAGLCODTAOSYTFG
1		1	1			CGHELDEEGLYCNSCLAOOCIN
		1	l			IQDAFPVKRTSKYFSLDLTHDE
23515	53883	A	23650	182	423	RLSIMASSLNEDPEGSRITYVKG
23313	22003	l^	23030	102	723	DLFACPKT\DSLAHCISEDCRM
			l			GAGIAVLFKKKFGGVQELLNQ
	1	İ	1			HYKEKGLRTSQLYG
	L		L		l	III VEVOLVIOČE IO

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23516	53884	Α	23651	1	249	
23517	53885	Α	23652	435	494	
23518	53886	Α	23653	163	578	RVTWLKKHLNSGNSDSVCMAS SLNEDPEGSRITYVKGS\LFEGP KTDFLAPCISEDCRMGAGIT\VL FKKKP\GGVQELLNQQKKS\GE V\AVLKRRWGRLYIFTWITKRK
						RAFAQSPLYENLTEEFRGQLKS HLSEEWSH
23519	53887	A	23654	1	2238	HL3EE W3H
23520	53888	A	23655	26	1941	RDAGTKWQRLGHLSCHCRKQA AREIINMHFQAFWLCLGLLFISI NAEFMDDDVETEDFEENSEEID VNESELSSEIIYKTPQPIGEVYFA ETFDSGRLAGWVLSKAK\KVD
						MDEEISIY DORWEIEEL KENQV PGDRGLVLKSRAKHHAIS AVLA KPFIFADKPLIVQVEVNFQDGID CGGAYIKLLADTDOLILENFYO KTSYIIMFGPDKCGEDYKLHFIF RHKHFKTGVFEEKHAKPPDVD LKKFFTDRKTHLYTLYMNPDD TFEVLVDQTVNKGSLLEDVV PPIKPPKEIEDPNDKKPEEWDER AKIPDSAVKPEDWDESEPAQI EDSSVVKPAGVLDDEPKFIPDP NAEKPDDWNEDTDGEWEAIPQ LINPACRIGCGEWKPPMIDNEY YKGVWIPPLVDNPNYQGIWSPR KIPNPDYFEDDHIPILTSTSALG
	53889	A	23656	297	398	ADHWAADGWRWKIMIANANK PGYLKQLMAAAEGHPWLWLI) LVTAGVPIALITSICWPTKVKDI RHSDLEYIPDICIPQTKGVLEQE EKVEKAALEKPMDLEEEKKQN DGEMLEKEESSEPEKSEEIEII EQQEESNQSNKSGSEDEMKEA DESTGSGDGPIKSLEKRRYRKD
23521 23522	53890	A	23657	286	430	TAGARCSSRTSSPGSCFNCOKS
23522	53890	A	23658	405	524	GHWAR\NARSPGFLLSRVPSVN WIHI ARMSVACLPGLFFICLRGDFKQ
		1				NLTAYCWKCYQL*RSLGTHS
23524	53892	Α	23659	ı	657	
23525	53893	A	23660	22	334	ALKHFCLCSLIFSVTTMKFLAV LVLLGVSIFLVSAQNP\TTAAPA DTVSSLLVLLMMKPLDAETTA AATTATTAAPTTATTAASTTAR KDIPVLPKWVGDLPNR

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nuelcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	eodon for peptide sequence	of peptide sequen ce	deletion, \=possible nucleotide insertion)
23526	53894	A	23661	I	270	
23527	53895	Α	23662	1	284	
23528	53896	Α	23663	111	281	
23529	53897	Α	23664	229	363	
23530	53898	A	23665	250	586	PVRHGAFFQDKSSDKKVQTKG
				[·		KRGAKGKQA\EVAN\QETKEDL
Į.						PAENGETKTEESP\ASDEAGEKE
	1					COCLINNHIPCLISGPCLPSCTIO
		1				RNIFINYFVNASFLVALETFLKK
23531	53899	A	23666	3	478	APAVGLSSSGGGRSGSGKAARF
23331	33677	l^	23000	ľ	770	REGCRRAAARMQQAPGTRLPR
ŀ		1				RHPARAFPAATMPKRKVSSAE
	i	1				GAAKEEPKRRSARLST\KLPAK
		1				V\EAKPKKAAAKDKSSDKKV\Q
		1				TKGKRGAKGKQA\EVANQETK\
	ł	l				EDLP\AENGETERLEEESSLLIEA
		1				GEKEKPKS
23532	53900	A	23667	1	252	GERERINS
23533	53901	A	23668	328	414	
23534	53902	A	23669	1	3156	
23535	53903	A	23670	1887	2036	
23536	53904	A	23671	3	212	
23537	53905	A	23672	22	539	YEGIVMKPLIIVNPADCIGC\RT
23337	35705	ľ.,	23072	[T		CE\VACVVAHPSEQELNADVFL
1	i					PRLKVQPIWTAVSAAR*CCPFSS
						ENAP\CVGA\CPVGA\LTMGEQV
1					i	VQTNSARCIGCQSCVSACPFGM
}						ITIQSLPGDTRQQIVKCDLCEQR
1						EEGPACVESCPTOALOLLTERK
						LRRVRQQRIVVSGENPL
23538	53906	A	23673	135	302	PIHMKTROPRDOFSADCVFLOS
				1		TCPVPC*RNRL*TPGGYWRTRR
						DYLRCDRSVHV
23539	53907	A	23674	175	794	LSTHSQPIADAADPPPHLHTGEL
				1		LCSAHOADFAVSAEPARPANN
				1		NAATTAPVHAVAITPPAYPDCF
						RTLPKSLSHODSTORLYGIARL
						LLRISTSGPPPSRLKVORLDSISA
						PV/MCHOCENAPCVGACP/VGA
						LTMGEQVVQTNSARC/IGCQSC
						VSACPFGMITI/OSLPGDTROOI
l						VKCDLCGALRDCVAVLRORRN
						SRVVLDACTRA
23540	53908	Α	23675	2	434	LISLPFPPDSLAHRPPSSAPTEPQ
	1	Γ.		Γ	1	GLFTVAAPTPGAPSPPATLAHL
l	1	1	1	1		LPTPAMYSLLETELKNPVGTPT
I			1			QAAGTGGPAAPG/GRKMALEN
l						PKMHNSEISKRLGADWKLLTD
1	1		1	1		AEKRPFIDEAKRLRAVHMKEYP
1				1		DYKYRPRRKTKTL
23541	53909	A	23676	3	264	DIKIN MICHEL
1-7741	123707	LC.	22070	Ľ	,	

SEQ ID	SEQ ID NO:					Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23542	53910	A	23677	1	2377	MDDTFLNYKYKNKCGPKHIKK
						KEEEEEEDNNSNDSKNKDKTL1
		1	İ			TGSSGSAKIFLMRGGRRQLLLP
		l	i		l	TLSLCLLSVQRSPGSHESGPPQS
		1				MNPRLSPQGKYLKNDFLDVIG
		1				MCDIACHLVVELLVGNNGNFL
ļ.		1				AHAFAHVEVVAQQNQRENAA
			Ì			RPNGGSVMIGQGASLRTCQSRV
		1				LRVARGGPSPNPGESEQGLFTV
						AAPAPGAPSPPATLAHLLPAPA
						MYSLLETELKNPVGTPTQAAGT
			l			GGPAAPGGAGKSSANAAGGAN
		1				SGGGSSGGASGGGGGTD\QDR
					1	VKRP\MNAFMVWSRGQRRKM
		l				APENPKMHNSEISKRLGADWK
		1				LLTDAEKRPFIDEAKRLRAVHM
1						KEYPDYKYRPRRKTKTLLKKD
1		1				KYSLPSGLLPPGAAAAAAAAA
1				1	1	AAAAAASSPVGVGQRLDTYTH
						VNGWANGAYSLVQEQLGYAQ
į.						PPSMSSPPPPPALPPMHRYDMA
						GLQYSPMMPPGAQSYMNVAA
İ			İ			AAAAASGYGGMAPSATAAAA
						AAYGQQPATAAAAAAAAAAAM
			ŀ			SLGPMGSVVKSEPSSPPPAIASH
				l	1	SQRACLGDLRDMISITGRENIFV
		l			1	VGDPAACLLPIEMEETFLPWPL
		1		1		GFEDRLHLEGPRPVQEIPSFLRK
		1	l			VPNCGQWLLQTPSVEPADPWE
		1	1			NPQAFPPSLQQEQPCLSLIVGIF
		1		ĺ		VEAFGGRGRWGSKGKKPKRYR
		1				TVEWMRIHQPSICCLQETHLTR
23543	53911	A	23678	1	399	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \-possible nucleotide insertion)
				sequence		
23544	53912	A	23679	13	1216	LGNKRAGLRRADGEAPSALPY
						CYFLQKDAEAPWLSKPAYDSA
	1					ECRHHAAEALRVAWCLEAASL
						SHRPGPRSGLSVSSLRPTRKMA
	1					TTFLAHEKIWFDKFKYDDAERR
						FYEOMNGPVAGASROENGA\T
						VILRDIARARENIHKSLAGSSGP\
						GAS\SGTSGDHGELVVRIASLEV
						ENQSLRGVVQELQQAISKLEAR
						LNVLGKELRLAHRGTAPOTOH
						VSPMR\QVEPP\AKKPATPAEDD
			l			EDDDID\LVGRDNE\EEDKEAA
						OL\REER\LR\HYAEKKAOESLP
	i					LGGQVPPSLLDVK\PWDDESDM
						AQLEACVRSIQ/LGTGRSWGAS
			l			OMVAVGYG\IRKLOIOCVVEDD
	İ					KVGTDLLEEEITKFEEARARSV
						DIASFSTRSEALKCVYVAHVRE
						GPCTRSKDLRPAKKKKKK
23545	53913	Α	23680	1	2066	MAAALRAPTQVSAASSRPPLPS
						PPETEGRVKGFLQPGPHCPAQC
						LCSGDDGSNEDLEQKREVIFPR
						PHEAPVAEHRQQTVGLRELVAI
						PKRKEETTAIVQVFVAFEDVAI
						YFSQEEWELLDEMQRLLYRDV
						MLENFAVMASLGCWCGAVDE
	l					GTPSAESVSVEELSQGRTPKAD
			İ			TSTDKSHPCEICTPVLRDILQMI
	1					ELHASPCGQKLYLGGASRDFW
	İ					MSSNLHQLQKLDNGEKLFKVD
				ı		GDQASFMMNCRFHVSGKPFTF
1	l			1		GEVGRDFSATSGLLQHQVTPTI
			l			ERPHSRIRHLRVPTGRKPLKYT
						ESRKSFREKSVFIQHQRADSGE
						RPYKCSECGKSFSQSSGFLRHR
	ŀ		1			KAHGRTRTHECSECG\KSFSRK
	l	1				THLTQHQRVHTGERPYDCSEC
						GKSFRQVSVLIQHQRVHTGERP
	[l	İ	l		YECSECGKSFSHSTNLYRHRSA
	İ	l				HTSTRPYECSECGKSFSHSTNLF
						RHWRVHTGVRPYECSECGKAF
		l				SCNIYLIHHQRFHTGERPYVCSE
	1	l				CGKSFGQKSVLIQHQRVHTGER
	1	l				PYECSECGKVFSQSSGLFRHRR
		l				AHTKTKPYECSECEKSFSCKTD
		l		1		LIRHQTVHTGERPYECSVCGKS
		l		I		FIRKTHLIRHQTVHTNERPYEC
				l		DECGKSYSQSSALLQHRRVHT
		l				GERPYECRECGKSFTRKNHLIQ
	1	ı		l		HKTVHTGERPYECSECGKSFSQ

SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO.	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \-possible nucleotide insertion)
23546	53914	Α	23681	Ī	468	MAAQVGAVRVVRAVAAHEEP
	ì	1				DKEGKEKPHAGVSPRGVKRQR
						RSSSGGSQEKRGRPSQEPPLAPP
l						HRRRSQPPTSWAAAVNDM/PAP
1						TVPGPVEPLLLPPPPPPSLAPAG
	1					PADDAPLPAPSTSGLFTFSPLTA
!	1					SAAGPKHKGTQGAAQAPSPPRP
						RW
23547	53915	Α	23682	950	1326	RPESQRANGVDSGPNLKTVPQP
	1	i				DTRKGSVLKWISKRGKPLAVEI
l			l			EESHCL\CLPLRTECLGIKP\IVH
					1	LFSCTRPVIVPSLELNYDVDSIA
l						HMFVADLLLMITLPSYDIPFYC
		<u> </u>				LVFQNLLVLEFQYL
23548	53916	A	23683	271	918	NQGLRDLGLCRICLVNKMFTSS
i						ILGKGHRHSLVSIKQEHSALRK
	l	1			1	AAG\PLPLKVGYC\QGFSPCDSL
	1	l				KYG\SWDEKDLTVPQPDTHKGS
1		l		1		VLRWISKRGKPLAVEIEEG\HCL
		I				PELPPGELECPG/ILKHGLYHWS
		l				SSEMGE\NRPMVG\ARHVYSNA
		ı				ALLFFTALRIFGQRRNINLGLRA
	1	i				HPG\IVPPLELNYDIDSSAHM/FF
20540	53917	١.	23684	,	942	LADLLLIITLPSYYIPFC
23549	53917	A	23685	346	1753	TWIPLSRTGRRFPVFSCSSSRYV
23330	33916	ľ	25005	370	1755	AAISFHKRWKEGRCFGRTRGE
1			İ			MGWEGRSRSLKELNTRNDWNP
		l		ì		GERELDFWRKITRFKEAASWTL
		1				EALEEAGSTARRRVWLEDGRR
i		1	1	Ì		TCEILSVLCTSSLSSNCASLDCV
		1	İ	Į.		TYKALSTTRLYDSVYWNKKPE
	İ		1	ļ		RSSSPATEOSWTENDFDKLREE
		1		l		GFRRI\NYSELKEEARTHGKEVI
				l		NLEKK\LDERLTRITNAEKSLKD
		ı	1			LMELKTMARELRDECTSFSSQF
		1				DPLEERVSVMEDQMNEMK\QE
		ı				EKFREKRIKRN/EKQSLQEIWDY
1		ı				VKRPNLHLIGVPESDRENGTKL
		İ		1		ENTLQEIIQENFPNLARQANIQI
				1		QEIQRTPQRYSSRGATPRHIIVR
1					1	FTKVEMKEKMLRAAREKGRVT
1		1		1		HKGKPIRLTADLLAETLQARRE
		1		1		WGPIFNILKEKNFQPRIAYPAKL
				1		SFISEGEIQYFADKQMLRDFVTT
		1		1		RPALKELLKEALNMERNNWYQ
	1	1		1		PLQKHAKW
-						
23551	53919	A	23686	73	327	GALQPTAALWEPLSGLAKAAA
23551	53919	A	23686	73	327	RSLSLQGGVEGDARAGTGAAR
23551	53919	A	23686	73	327	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23552	53920	A	23687	322	402	DQEPTNSGHIGNLEQCLGHGYQ R*LFP
23553	53921	В	23688	545	3063	
23554	53922	Α	23689		579	MKHEKRDRCAELTAILLNEQAS PMCTIVLGYTGTSTITLPRTSCH FAPRLSNSQUESSYQVGHKPES YCVRNWWVLGLTDFKNEAAD PHNSGQQLASPSGSRTGAAGG\ VPASPAPCARTPOPLGGRWAD APGTGGGIRRGGSGCTGTHGG GGRLRHGGLQSGGLPRGKAAK ARREJESSAGYSNSSSKCSII
23555	53923	A	23690	399	592	KCIHFPGPTPKFVCGVIVVTAVP /TFGM*TVYV**LPFSILAQNVG *RDDCDPGWKLNLFYAKKKK
23556	53924	A	23691	3	268	DRV/RPLRKNFQRNDQTAAFTV HENPLLCSYRCWYPGKQEQKK IKPQRGGKNRAEKLETLKSRAP LLFQRNAVPHQQRNKAGWRM TLTS
23557	53925	А	23692		484	CSFFLVSMVFTFWHDFAAAGTS CSFPCLVLSFRSSFRAGLVVTKS LSICLSVKYYFISSSLMKLSLAGY EIVG*KFFSLRMLNIGPHSLLAC RVSAERSAVSLMGFSLRVTRPF SLAALNIFSFISTLVNLTIMCLG VALLEKYLCGILCVS

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion. \=possible nucleotide insertion)
23558	53926	A	23693	182	2076	MQEWLRLTARKTELSRDSRSC GQLKTSLLISLLVESTOGRI+SF LVWTOTHSGGDGWRVAKAW RFKYHDTQLGGEKEEPQGHHA SDCRWRAQPHCRTLSAAVVNV AGFTVTFELGCLTLMKMREKD HHLCDNDIVLEPPDTAFPLDCL LSCHISTVHGLLRWNAIQYLAS SESSHPERSEGTPDRNOHPAL SLASFNGAQODRRTVQSLGEKP GLPEEMAAFMAGTEKEQNELG IEFFQLIPASKRTHSENALFSHS GDDTECSRSQQCSPPSYSRGPP RTFILATOPLCSKEAGTSPHEQ PTRTEKNRGSHLYHKSPCDHAS PGMSVTMSGMQQALPRQLTIL THYPPAFRSIKFIRMILKTEMLH THARSQEALLKDSSKTMLHSTL PSCVRKSNYHHRERKIKTEMAL FAKHRSNYAMFREKHIKTEAF PAKHRSRNGGMKVEVAPLITI SSDPLAEFILLPVTVTLCSASLEW LAPEGGMLSUNDERSTENEW LLPEGGMGATMENEW LLPEGGMGATMENEW LLPEGGMGATMENEW LLPEGHGGLLIPLSQQVKRV
						PVLVGVIDLDYQDEIDYYSSMK ERSSPATEQSWMENDFDELRE EGFRRSNYSELREEIQTKGKEV ENFEKNLEECITRITNTEKCLKE LMELKTKVRELREECRSLRSRC DQLEERVSAMEDEMNEMK
23559	53927	Α	23694	933	1118	RVERRRLQTIKLLRATGGNSNQ RQRS*KL*KKFRRMYN*NNQY REVLKGADGAESQGSRTT
23560	53928	A	23695	2	250	WCHTKLDGLSLFGESVALEEA GGQTALWLMRFLSRKANYVD KITTREVGHRE*LMSSRLHSR* WAASSVCSSSCVLLPSSHS
23561	53929	A	23696	1017	1230	
23562	53930	В	23697	62	1069	
23563	53931	A	23698	216	582	ILVPTLFWLGEFLLRDPLLV+W ASI*G*PD/AFSLAALNIFSFISTL/ RESDNYVSWSCSSRGVSL*RSL YFLNLNVGLPCQIGEVLLDNIL QSVFQLGSILPVTFGYTNQMQI WFHFIHFIFHH
23564	53932	A	23699	434	681	TLACLARLGKFSWIISCRVFSNL VPFSLSLSGTPIRRRFGLFT*SHI SWRLCSFLFILFSLNFPSRFISFV SSSITDTLSSS

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23565	53933	A	23700	290	1231	RSFRAIGEVGSIRGWGGGYAAO
1		1		I		LQSPSRLPVLGGGKKGGSRGV
						GOGSSRAEVGLGSQRERSGTCG
1						AEAAAAAAQLIAGTGSSLPYQG
1		1				PGONOTAIAPTCAPLSPCKAAA
		1				VIMGNIFGNLLKSLIGKKEMRIL
		1				MVGLDAAGKTTILYKLKLGEIV
İ		1				TTIPTIGFNVETVEYKNISFTVW
		1				GCGVARDK\IRPL\WRH\YFON\T
		ļ				QGFDILWSDSN\DR\ERVNEA\R
		1				EEL\MKNAGAETSLRDAVLPFL/
		İ				FGNKTDLRNAMNAAG\IQDKL
		l				G\LHSLRHRNWYI\QATCATRG
						NGLYEGLHWLANSVPKQEVES
						RTALTKHPNRP
23566	53934	С	23701	164	208	
23567	53935	С	23702	1	381	
23568	53936	С	23703	43	276	
23569	53937	С	23704	1	462	
23570	53938	С	23705	38	121	
23571	53939	A	23706	77	202	QKNYTQRPLPLHGLLELEVTCR CWSWQPLLEPLELRP\RLSD
23572	53940	Α	23707	123	209	
23573	53941	С	23708	1	411	
23574	53942	Α	23709	1400	1809	SSNPLSRPPLPTPVTPCPLLRLSL
		1				LRACCCCLFLPLRFFSCPLLCFL
			ĺ			LCFFFSVCSPAHKNFKMPKPQW
		1				SGIPTGPPPSGSCFKCQKSGHW
ì		1				A\RNACSPGFLLSHVPSVWDPT
		1				GNQTVQLAWQPLLEPLEL*P\R
		١.		_		LSD
23575	53943	В	23710	76	358	
23576	53944	В	23711	1	606	
23577	53945	С	23712	272	388	
23578	53946	Α	23713	1	259	PGVRGAQGGPSIPRQCEESAIGP
						KFAWIISSFHWKQGMFNMEAP
		1				SSLFFVNMCAVKKQTTWGRPD
		_				RGFIWQPLPEPLEL*P\RLSD
23579	53947	Α	23714	684	917	GDFGLRQWGFLNIQSCHLQTGT
		1				I*LPLFLIEYPLFLSPA*LPWPELP
		1				TLCCIGVVREGISVLCQFSKGM
		<u> </u>				LPVFAHSV
23580	53948	A	23715	618	683	
23581	53949	A	23716	306	710	DRVSLLTPRLGVQWPDPGSLQP
	1	1				RPPGFKRFYCLSLPSS*DYRHEP
		I		L	<u></u>	LLPAYFFPYRIIGWSRL
23582	53950	Α	23717	2	88	

SEQ ID NO:	of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	location of first eodon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop coden, /=possible nucleotide deletion, \:=possible nucleotide insertion)
23583	53951	A	23718	972	1967	FPPGRGROPGRAGCSRGRARA CPKPVDHVKRPMNAFMVWSR AQRRKMAQENP;QMHNSEISKR LGAEWKLLTESEKRPPIDEAKR ILGAEWKLLTESEKRPPIDEAKR ILLKKDKFAFPVPYGLGGVAD AEHPALKAGAGLHAGAGGCL PSELLANPEKAAAAAAAAGA VFPPQSAAAAAAAAAAAG YPSPQSAAAAAAAAAAAGS PYSLLDLGSKMAEISSSSGLPV LMMPMMHINSLGKNLALNLFV YNDANSMLDLCYDGVDCKGPI CFCATGGYEDIFKAGYGSVVD YYNDGSDLTTPLEIATVIPTFSN HLMSQRSTWRQNPPLAKRLP LITEDSDGDDH
23584	53952	A	23719	2	144	RPQRRP/LPPPTGSPLKHPGGAS GAAGAPIGEPQHPGAVVSVRA WSLCS
23585	53953	A	23720	219	897	PGRAGGTLGLOSRRDOTREKG VEETPHRLHARTDTTAPGCCGS PIGAPAAPEAPPGCFEGPRWVE GVKKASPPLPAELIGWLVNFPL ARPVAPPSLSOPPRRPPEGVCG SWPGRSEGERGKAKNTAESTPR QNLHRKYTIIVTHYGRNRKGQ NISFSLVCEFIPSLADWTTVAL LTWTSPKDNSLTEESWLREGN VHKGMDEYVRAQSFLILMTLE NSGGSVVSK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
ł	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23586	53954	Α	23721	937	2730	ТИРНСИДНҮҮНИННЕНСИНСИ
					ŀ	нининсининунининской
	ĺ		ŀ			нүннинисинсинининсин
				ì	ŀ	ННҮНННҮНННСНДННҮНННН
	1		l			НСИДИНУНУНИНИСИДИНУН
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ļ						Y НИНИНСИDНИЧНИНИНСИD
	l			i		ннүнининсноннүнүнинс
						НННКҮНННННСННСННННН/Р
1			l	i		нинининисниннунининс
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						ннинүүнинсинсининүн
						ннинсирннунинсинсини
						НҮНИННҮНДНИҮНИНННСИ
					•	НСНИННИННОННИННЕНСИН
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1		i				ннисинсунинуниннунси
1		1				ННҮННННҮНРҮНҮНОНННН
Į.		l				ҮНРННҮНННҮҮНСННННОҮН
						НРННСНННННҮНОННҮҮРІТІІ
						TTTITTITTIIITVTTTTFTTTTNT
		l				TAPTTTPVTAITTTTTTTTTTTTT
		1				TPITITTITNITTTTVTTTVIITTTF
		l				ТТТІНРҮННННННРУТІТТІТ
		l	İ			TATHTTTTSITSITTHTTTIVTTH
1		1				ТТТААТНИНҮНРҮҮННҮННН
1		1				ННННТНЅҮҮНҮНННННННСГ
						НҮНИНҮНИН
23587	53955	A	23722	2348	2571	
23588	53956	A	23723	4919	5142	
23589	53957	A	23724	2	370	
23590	53958	A_	23725	588	701	
23591	53959	A	23726	295	521	
23592	53960	A	23727	2837	3061	
23593	53961	A	23728	272	436	
23594	53962	A	23729	239	514	TOP DOCUMENT OF STREET
23595	53963	Α	23730	105	367	TGRPPELLLHLRSLSPPLL/CSSP/ TPPPQLPPPPVSLASGYPLLHPL
		1	1		l	ARYPACPLPPRRARTEORGGEO
						PGVGASPLPPRRAKTEQRGGEQ
22.506	52064		22721	999	1146	PG V GASPLPIPGK WPG IEA I
23596	53964	A	23731	1084	1145	
23597	53965	A	23132	1084	1893	l

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \mpossible nucleotide insertion)
				sequence		
23598	53966	A	23733	1605	2296	HSIHSSVDHPENRIYRRKIEELS
23376	33700	*	23,33	1005		KRFTAIRKTKGDGNCFYRAL\AI
1	1					PYLESLLGKSREIFKFKERVLOT
				1		PNDLLAAGFEEHKFRNFFNALR
					· ·	AGRAOFYSVVELVEKDGSVSSL
						LKVFNDP\GASDHIVQFLR/LCF
			1			TSAFIRNRADFFRHFIDEEMDIK
			ł			DFCTHEVEPMATECDHIOMTA
						LLQALSIALQVEYVD\GWIPALT
						OHVFLEAATPSVYMLYKTSOY
	l					NILYAADKH
23599	53967	A	23734	529	1830	AGYGDRTMLARKSIIPEEYVLA
23399	33707	<u>۱</u> ^	23/34	329	1050	RIAAENLRKPRIRDRLPKARFIA
			1			KSGACNLAHKNIREOGRFLODI
		1				FTTLVDLKWRHTLVIFTMSFLC
			- 3			SWLLFAIMWWLVAFAHGDIYA
						YMEKSGMEKSGLESTVCVTNV
		l				
		1				RSFTSAFLFSIEVQVTIGFGGRM
						MTEECPLAITVLILQNIVGLIINA
						VMLGCIFMKTAQAHRRAETLIF
			l			SRHAVIAVRNGKLCFMFRVGD
						LRKSMIISASVRIQVVKKTTTPE
						GEVVPIHQLDIPVDNPIESNNIFL
		1		}		VAPLIICHVIDKRSPLYDISATDL
		l				ANQDLEVIVILEGVVETTGITTQ
		l				ARTSYIAEEIQWGHRFVSIVTEE
		l				EGVYSVDYSKFGNTVKVAAPR
		l				CSARELDEKPLPSLFRPSQKSEL
		l	l			SHQNS\LRKRNS\HRRNNFHRRN
		l				N\SIRRNN\SSLMVPKVQFMTPE
		_				GNQNTSES
23600	53968	A	23735	968	1243	author is the state of the stat
23601	53969	Α	23736	541	1010	SVTSCLKVTSVCLTSAAASCRH
						RRSVEEYLVS*KLPM/WSRRRK
	l	1	1	l		AAALTES/V/CIVFWLASWMMS
	1	1		l		RWLHYPDWVTIIAQQVLMNVV
	l	l	l	l		SVMQADCRMNLLEIFLRWLMR
1	1	1		l		KIFHNILRHFSQLLNVPITRPFS
		_				WILRPF
23602	53970	Α	23737	766	1910	LLTARLTWVSARLMSYVLLMC
	l	1		l		*LFPRLLSCLTTPPHCSFSICFVI
		I	l	l	l	CSRTLILKGSSLMYVFCLPNTAI
		I	1	l		GGSSRVDRCRAATSRGSSRVDL
		I	l	l	l	TL*KYSLVGPSVPLVSSV**SDW
		1		l		PLKLLLMTMMS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23603	53971	IA	23738	581	1434	I KLPMGVAVVKLLHLPKVIIVFW
						LASWMMSRWLHYPDWVTIIAQ
						OVLMNVVSVMOADCRMNLLEI
		ł				FLRWLMRRASDICSSIHGD*EPL
		1				L/YAVFGKQNTYIRLEPFKINVL
i		1				EQITKHIEKLQCGGVVKQLSRR
ļ.	ŀ	l			i	GNNOHISSTYDINRADTOVRRA
		1				VNNYDIIVMSNSFNGQSEHQV
1						WIDRKAGECLLHEYEDLVPIRD
1					i	TLRLFPGGRYLPRAKHVAPSEP
		ı			l	DPEQDEQKLRFCKRHLYGKOP
1		1				RSPVERVWDHGPTRIVGLIISLG
		i	ľ	1		PRSHSYRRSDNQTGTTVPLVSS
23604	53972	A	23739	515	774	LFPRLLSCLTTPPHCSFSICFVIC
23004	337/2	l^	23139	1313	l''*	SRTLILKGSSLMYVFCLPNTAIS
		1		1		GTTVPLVSSV*LLVWDHGPTRI
		1		i		VGLIIRLGPRSHSYRRS
23605	53973	A	23740	322	763	CRMC*LFPRLLSCLTTPPHCSFSI
23605	339/3	A	23740	322	163	CFVICSRTLILKGSSLMYVFCLP
1						NTAIVMALSPRGWRSKFGMPV
1		1				EDTSLSTPAAPMVDSLIARVGV
[1				MARGNAITLPVCGRDVKSTLE
l		1				VLRGDSVEKTSRVWSGNERDO
		ı				ELLTEDALDDL
20.606	53974	+.	23741	11	127	MKLMETLNOCINAGHEMTKAI
23606	539/4	A	23/41	11	127	AIAQFNDDSPEARKITRRWRIG
		1		1		EAADLVGVSSQAIRDAEK*RRL
		ı				SPLHSLMMTARKRGK*PGAGE
23607	53975	A	23742	9	437	WLGLGNWKERLGTOTRMIWS
23007	33913	l^	23/42	ľ	437	OKTARRLCRTFSGIPAYSMCI/CI
						GMAERMWCDRNRERHTVSSSG
		1		1		GNRLPNPDWAPGPPRIVGLNIS
						LGPTVPTRIVGLIISLGPRSHSYR
						RSDNQTGTTVPTRIVGLILVLGP
						WSHSYRRSDY
23608	53976	١.	23743	766	767	LLTARLTWVSARLMSYVLLMC
23608	539/6	Α	23/43	/66	/6/	
l		1		1		*LFPRLLSCLTTPPHCSFSICFVI
1					1	CSRTLILKGSSLMYVFCLPNTAI
	1	1				GGSSRVDRCRAATSRGSSRVDL
				1		TL*KYSLVGPSVPLVSSV**SDW
		_				PLKLLLMTMMS
23609	53977	A	23744	573	1085	VCHQILTHLPALLSGARMMWD
						QLWRGRP*RRCQQPQCGG*WH
1		1			1	CFWATAHHSPLDSTLNHLHCLP
						RGP
23610	53978	Α	23745	2	379	WLTRSSCSWRLPPRRKKKGAL
		1			1	V*PVWLPPGGVAVLLQLGCCV
		1		1	1	LQDVSCKRLCYNRKGQSEIVL
		1		1	1	MS*ITMWKLFPRPPLLPFSTCIC
		1		1	1	*KTCGKRGRSK*KVDILVLSIRF
		1		1		SAIKLEPLDCHENAWL

SEQ ID	ISEO ID NO.	Inter-	SEQ ID NO:	Nucleotide	Nucleotide location of loca	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		丄			L	
23611	53979	Α	23746	575	576	TGIDSDQLNGRDWRRALSVSPA
						PDSGFPRCNGDEQAGI*LLGTG
				1		WAEKCHRLAFPPAGSTAGR
23612	53980	Α	23747	1	723	MKLMETLNQCINAGHEMTKAI
		i .		1		AIAQFNDDSPEARKITRRWRIG
		1				EAADLVGVSSQAIRDAEKAGR
		1		1		LPHPDMEIRGRVEQRVGYTIEQI
		1	1	1		NHMRDVFGTRLRRAEDVFPPVI
		1				GVAAHKGGVYKTSVSVHLAQ
		1				DLALKGLRVCSWKERPPGNSL
		1				NVSRMVWDHDPTRVVGLIIGL
		i	1			GPRSHLYCRSDYQQRS\TSVCG
		1		1		CMPAVDCCCVLLIHNILRTVM
		1	Ì			WTKYLVTQAVPAR*PGCIRCKC
23613	53981	A	23748	.1	1101	
23614	53982	Α	23749	5	205	YCSSLSFTAAITSRAFLPDRRITT
		1		1		PPATSPSPFSSAMPRRISGPIWTF
		1				ATSRR*TGTPFSLAFKTM
23615	53983	Α	23750	I	761	
23616	53984	В	23751	56	2188	
23617	53985	Α	23752	627	1272	
23618	53986	A	23753	448	2218	VRRQRSDPERSDARMVRFCNL
		1		1		YI*RKNPFILH*LFR*TLRQTKPD
		1		i		NSAGKCVKI
23619	53987	Α	23754	17I	959	AASRGPDAPGLEMAGNCSWEA
		1				HPGNRNKMCPGLSEAPELYRR
		1				GFLTIEQIAMLPPPAVMNYIFLL
i		1				LCLCGLVGNGLVLWFFGFSIKR
	İ	1				NPFSIYFLHLASADVGYLFSKA
		1				LFSILNTGGFMGTFADYIRSVCR
		1		1		VLGLCMLLTG\VLLMPAGWDE
		1				RGVRSERAWCTVETGRVHRSR
		1		ł		VRVSRNTARDPANGLGHLLQS
		1		İ		LGMPALEPGPTLGSALRWEFLQ
		1		1		TASPQLPVQPLHPAAATPQRSL
		L				RGRGRGPKARALGLGPRSTSYR
23620	53988	Α	23755	664	838	
2362 I	53989	Α	23756	2732	3007	
23622	53990	Α	23757	1522	1706	CHYFACMWSGCEVYS*SAPG**
		1		1	1	C*EDLSGLVRCRCLHSNAETPI*
		Ц.				FAANADTSPRGPFLCR
23623	53991	Α	23758	537	795	ADCPLHMGRPCLID*GPCRTE*P
		1				RCAVVCLLWIAAVSCLSTT/SLR
1		1				TGI/DGPKYLVTQAVPARYPGCI
		1			į .	RCKCVAVDELASSDMRLPCS
23624	53992	Α	23759	2074	2276	
23625	53993	Α	23760	36	206	ECWIWCELGAELHHQP*ASSH
	i	1		1		WKSSYRQQSSLPRCRASQRTLH
	1	1		1		LLLQGWAVLGPLE
23626	53994	Α	23761	444	725	
23627	53995	В	23762	I	1212	
						

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, !=possible nucleotide deletion, !=possible nucleotide insertion)
23628	53996	A	23763	11	972	RFSKSLLDSGVQLASPSGSRTG ATGGAAHSPARAPALHSPWAV DGTRGRLAGTAGLWSPPAFH FKVSLSLPFSCVSSPPGPPLGPS PRKHGMADDAGAGVGGRRPF RGPMGSNGGGRGFRGFGSGIRHQ GCGCGQGQGGRGVRGGKAE YKEWMTVTKLGHLVKDIKIKS LEEIYLFSFKTSSSLKDEVLKI MLVQKGTHTSQCTRFAFVAT GDVNGHISLGKCSKEVATAIC GAIILAKLSIVPACRGYWGNKIS KPHTIPCKVAGCCGSVLVCLIS YPRGTGIVLAPCEGDRGLRSS PASLSPESCKDGSHSRSF
23629	53997	A	23764	2	260	LLEECRSLRSRCDQLEERVSAM EDEMNEMK*EGKFREKRISPKR APERSAKHGKEQPVPAAAKSC QNVNTIETRKKLHQLTSKITS
23630	53998	Α	23765	3	1150	
23631	53999	Α	23766	1	1521	
23632	54000	A	23767	1	753	
23633	54001	Α	23768	1	3351	
23634	54002	Α	23769	1	1740	
23635	54003	A	23770	1	1353	
23636	54004	Α	23771	1	1932	
23637	54005	Α	23772	1	1221	
23638	54006	A	23773	I	1214	MAKKQNRKTGNSKKQRASPPP KECSSSPATERWYENDFDELR EEGFRQSNYSELWEDIQTKGKE VENFEKTLEECITRITNTEKCLK ELMELKTKAGLEKECRSLRSR CDQQGERASVLEVLEVLARQL GRKRK*RVFN*EKRKSNCPCLQ TT*LV1*KTPLSQPKISLS**ATS AKSQDTKSMYKNHKHSYTFIT *ESNLQES*NWISSLHLIQKSIQD GLKT*TLDLKPSKP*KKT*ALPF *T*AWARTSCLKHQKQWQQRT KLTNWFLN*HSTLLYGRAN URQLQ*TOTNLQEKNKQPHQKV GEGHQQTILKRRHLCSQKTHE KMSITTGHQRNANQNHNEIPSH TSYNONH

SEQID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
22620	154007	+	23774	11157	2085	OCOMPOS A DECETTIVO DEPLACA
23639	54007	A	23114	1157	2085	QGRPTDFADFGTTIKQDFRLLG
		1			ļ	QTSVDRLLQLSQGQAVKGNQL
		1				LPVSLVKRKTTLAPNTQTASPR
					l	ALADSLIAAGTTGFPTGKRAVS
			ł			ATQLMTQFSPAREQSWRKN/DF
		1	l			DELRKE/GFARSNYS\EKCLKEL
	l					MELKTKARELREECRSLRSRCD
				l		QLEERVSAMEDEMNEMKREG
			1	1		KFREKRIKRNEQSLQEIWDYVK
						RPNLRLIGVPESDAENGTKLEN
			1	1		TLODIIOENFPNLAROANVOIOE
		1				IQRTSQRYSSRRATPRHIIVRFT
		1				KVEMKENMLRAAREKGWVTL
1		1		1		KGKPIRLTADLVAETLQARRE
		1				WGPIFEEP
23640	54008	Α	23775	1	1707	
23641	54009	Α	23776	1	1899	
23642	54010	Α	23777	1	2781	
23643	54011	Α	23778	1	831	
23644	54012	A	23779	1	3264	
23645	54013	Α	23780	1	882	
23646	54014	Α	23781	1	1853	
23647	54015	A	23782	1	550	
23648	54016	Α	23783	1	831	
23649	54017	В	23784	64	3436	
23650	54018	Α	23785	1	1422	
23651	54019	A	23786	1	773	MGKKQSRKTGNSKKQSATPPP
					1	KERSSSPATEQSWTENDFDELR
				ŀ		EEGFRRSNYSELQEETQTKGRE
			ļ		Į.	VENFEKNLDECITRTTNTEKCL
			İ		l	KELMELKAKAWELREECRSLR
						SRCNQLEERVSVMEDEMNEMK
		1				REGKFREKRIKRNEQSPQVIRD
		1	1			YVKRPNLRLIGVPESDGKNGTK
						LENTLQDIIQENFPNLARQANIQ
						IQEIPRTS/QRYSSEEHSIIGLIGR
		1				FPS*RTDMFRAPKKREOKRNNR
		1				KIQKEEGGRTAPTGI
23652	54020	Α	23787	ī	2775	
23653	54021	A	23788	1	2712	
23654	54022	Α	23789	1	1590	
23655	54023	Α	23790	I	547	MGKKQNRKTGNSKKQSASPPP
	1	1	1			KECSSSPATEQSWTENDFDELR
	1	1	ŀ			EEGFRRSNYSELREDIQTKGKE
	1	1				VENFEKNLEECITRITNTEKCLK
	1	1				ELMELKTKARELREECRSLSSR
1	1	1				CNQLEERVPAMEDEMNEMKE
l	1	1				D/GNRYDDRRQ/RKEEIMIKSW
!	1	1	l			RFCCRHLIRVVRGWGPSKYLWI
	İ	1	١.	1		TMVVTLAGNPO
23656	54024	A	23791	1	710	
	1	1				

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23657	54025	A	23792	ı	895	
23658	54026	Α	23793	1	1938	
23659	54027	Α	23794	1	1413	
23660	54028	А	23795	985	1219	LCSGSVERCTESYLHSSHCLVQ QAGVTGDEGGSCY*DPPSHHRS HPRKLKKLCEQEFSVTSAPWRK CRSLHGRFPGLP
23661	54029	Α	23796	I	3570	
23662	54030	A	23797	1	634	
23663	54031	A	23798	I	4551	
23664	54032	А	23799		4721	MHPSQRTPYQSRSPATIFATVST FKSHKSNLETKHPQQVPLSSLG LAANISSEDIESEAEDNGKWES VDRTSFFPALTSFQQSMIHKKI MTVFKSVEPEDGLGRPTLAFHE DRCQLSIQNSLDAGGGCPSMA APPAEVHELASGLISAAALLLSP AASGRVSKHQHIPTGFPDPFKA LHKGKLTTRKDIYTENPSVHHH HQRPKVDKTTKMGKKQNRKT GNSKTQSASPPPKECSSSPATEQ SWMENDFDELREE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
ĺ				sequence		
23665	54033	A	23800	1	2152	MDILGWASRAHKWKGRATGT
ĺ						HVFNLMREYLISLQCCASEQAA
ĺ						SPEHGASFRVWKSLGFQKLILS
ĺ						LICPQCLQSLTRETWMRFKSKK
ĺ		1	1			TKGPDLRLVMDANKWEFRRH
ĺ		1				ALRPALTFPNEGGDLGVEALPV
ĺ		1				ALPPEAHVGGVVLPAALVRGA
ĺ		1		1		HGEQLPAIPGPEEFALPAVQAR
ĺ	1	1				AARPERQVVVIVVNEAGVDPA
ĺ		1		1		GVEGLAPVPVLPGVRLEATAPK
ĺ		l				QPPRLPARLRAPLPSGRKLNAK
ĺ		1				CPRLRWGFFVQDGDGTELACL
ĺ		l				SEGDGGCCQEAPLDHRPRRLRG
ĺ		l				LAGSAGAREAREPRLRPGDWG
ĺ		1				NRIARVRAPEEAAAALSRPARV
ĺ		1				AAALAKTHRREHRARSQKLVM
ĺ		l				SSVQQGKLTTRKDIYTENPSVH
ĺ		l				HHHQRPKVDKTTKMGKKQNR
ĺ		l			İ	KTGNSKKQSAF\PPPKECSSSPA
1		l	ŀ			MEQTWT\DNDFDELREEGFRGS
ĺ		l				NYSELREDIQTKGKEVENFEKD
ĺ		l				LEECITRITNTEKCLKERMELKT
ĺ		l				KARELREECRSLRSRCDQLEER
ĺ		1				VSAMEDEMNEMKREGKFREK
ĺ		1		1		RIKRNEQSLQEIWDYVKRPNLR
ĺ		1				LIGVPESDVENGTKLENTLQDII
ĺ			İ			QENFPNLARQANVQIQEIERMP
ĺ		1				QRYSSRRATPRHIIVRFTKVEM
		l		1		KEKMLRAAREKGRVTLKGKPI
		1		1		RLTAVLSAETLQARREWGPIFN
				1		ILKEKNFQPRISYPAKLSFISEGE
		1		l		IKYFIEKQMLRDFVTTRPALKE
23666	54034	Α	23801	1	2109	

SEQ ID NO:	of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
23667	54035	A	23802		2544	MSVHSWKPSWIRPIRSLSPKRRI GTOGTSEVPDTHICOTKERISLSPKRRI GTOGTSEVPDTHICOTKERISLS SGPLLLSVHPVGKLSAVVGRVF HIILSOGRERRRRAVIASRPSEVPG SSHQCVPDSGCRSSVGAHPVREP KQGEALPHSGSPREGKLTNRKD HITKNPSVHIHHHORPKVDKT KIGKKQNRKTGNSKTQSASPPP KECSSSPATEQSWMENDTDELE REGFRÇSNYSELRKDIQTKGKE VENFEKNLEEGKTRITNTEKCL KELMELKTKARELCEECKERS RCNQLEERVSVMEDEMNEM REGKFREKNIKRAELCEECKERS RCNQLEERVSVMEDEMNEM REGKFREKNIKRNEGSLQEIWD VYKRFNLELIGPYESDVENOTK LENTLQDIIQENFPNLARQANV VYKRFNLELIGPYSSRATPHHIV RFTKVEMKEKMLRAAREKDRS TRQKVNKDTQELNSALHQADL LIDIYRTLHPKSTEVTFSAPHHT YSKIDHIVGSKALLSKCKRTEII TYYLSDIISAIKLELRIKNLTYYQNI NYLLINSLLLNDYWYNNEM KABIKMFFETNENKILLELRIKNLTYYQNI KABIKMFFTNENKILLELRIKNLTYYQNI KABIKMFFTNENKILLELRIKNLTYYQNI KASRRQETKIRAELKEGETQ KTU,KINESRSWFFERINKIDR LARLIKKREKNQIDTIKNDKG UTTOPTEIQTITREYYKHLYAN KLENLEEMDTFLSFVITJERLN,PI

SEO ID	SEO ID NO:	Mct	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
			00000		Lone	NAME OF THE PARTY
23668	54036	Α	23803	1	1886	MKENFKHRHVVAFISSQEAQE
						DSREEEDRLESRFYFLFLLELSY
		1				KTTGLNREPTVSLNSKSGTPMD
		1		İ		QNEHSHWGPHAKGQCASRSEL
						RIILVGKTGTGKSAAGNSILRKQ
						AFESKLGSQTLTKTCSKSQGSW
						GNREIVIIDTPDMFSWKDHCEA
			l			LYKEVQRCYLLSAPGPHVLLLV
		ĺ				TQLGRYTSQDQQAAQRVKEIF
İ				l		GEDAMGHTIVLFTHKEDLNGG
		ł				SLMDYMHDSDNKALSKLVAA
						CGGRICAFNNRAEGSNQDDQV
						KELMDCIEDLLMEKNGDHYTN
1						GLYSLIQRSKCGPVGSDERVKE
		l			ľ	FKQSLIKYMETQRSYTALAEAN
		l				CLKGALIKTQLPKVDKTTKIGK
		l				KOSRKTGNSKKOSTSPPPKERS
						SSPAMEOSWTENDFDELREEGF
						RRSHYSELPEDIOTEGREVENFE
	Į ·					KSLEECITRITHTEKCLKELMEL
i						KTKARELREECRSLRSQCDQLE
						ERVSAMEDEMNEMKREGKFKE
1						K\RIKRNEQSLQEIWDYVKRPN
1						LRLIGVPESDAENGTKLENTLO
l						DRIOENFPNLAROANGFCRFRN
						HHOTGFSPAGANORGPLAATLS
						GPGGEGOSAVARLTGEKKNHP
1						GAQYANRLSPRVGRFINRSWH
						DRFPDWKAGTGLKSAIFAWE
23669	54037	A	23804	1	4134	Did 10 HE IS I SERSAH AWE
23670	54038	A	23805	1815	1849	
23070	134038	ΙΑ.	23003	1013	1047	

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for tast amino acid of peptide sequence	*-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
	sequence		09/540,217	sequence	or pepude sequence	detection, v-possible nucleotide insertion)
23671	54039	A	23806	1	4016	MGKKQNRKTGNS\KTQSAFP\P
		1				PKECSSSPATEQSWMENDFDKL
					1	REEGFRRSNYSELR\EDIQTKGK
		l				EVENFEK\NLEECIT\RITNTEKC
						LKELMELKTKGSRNYVKECR\S
		ŀ				LRSRCDQLEERVSAMEDEMNE
			1			MK*EGKFREKRIKRNEQSLQEI
						W\DYVK\RPNLRPIGVSESDGEN
						GTKLENTLQDIIPENFPNLARQA
						NVQIQEIQRTPQRYSSRRATPRH
					İ	ILVRFTKVEMKEKMLRAAREK
		l				GQVTLKGKPIRLTADLSAETLQ
		1		8		ARREWGPIFNILKEKNFQPRISY
						PAKLSFISEGEIKYFIDKQMLRD
		Ì				FVTTRPALKELLKESLNMERNN
ŀ		l			l	WTSKQIIQEDGYSEEEGWLCRA
ŀ		l				VIYSHTLQAIMAIVKATGNLQT
		l				DFANPFHTDDARQQFAMSYTV
		1				VGQGALPEDLSRCHLQTEAVQ
		1				RHPESLGCPLDLLHGMPVSSCF
						GYPQDTSSTV
23672	54040	Α	23807	2059	3153	
23673	54041	Α	23808	6	188	
23674	54042	Α	23809	3	1014	VACRPLPRRRAQPATFIPRVAV
		l				KMALVASVRVPARVLLRAGAR
		l				LPGAAFGRTERAAGGGDGA\RP
						\FGSQRVLVEPDAGAGVAVMK
İ		1		1		FKNPPVNSLSLEFLTELVIS\LEK
		l			1	LENDKSFRDVILTSD\RPGVFSA
			1		1	GLD\LTEMCG\RSPAHYAWVLE
						RPVQELWLRLYQSQL\VLVSAI
	1					NGA\CPAGG\CLVALTCDYTIL\
						AGNPRYCIGLNE\TQ\LGIIAPF\
l		1	1		1	WLKDNPWRTPIGHRAAESA\LQ
		l	1			LGLPLPPPAEALQVGIVDQ\VVP
					1	\EEQVQSTALSAIAQWMAIPKP
					1	MAHK\LTKAIMRKGQGPPAWS
1		1			1	RKGYADVQNF\VSFISKDSIQKS
	L	L	L			LQMYLERLKEEKG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23675	54043	A	23810	11	2665	MTTSIROFTSSSSIKGSSGLGGG
23073	34013	ľ`	23010	ľ	2000	SSRTSCQLSGGLGAGSCRPGSA
						GGLGSALGGSSYSSCYSFGSGG
						GYGSSSFEGVDGLLVGGEKAT
l		l				MONLNDRLASYLDKVRALEEA
				i		DTELEVKIRDWYOROAPGPAR
				i		DYSQYYRIIEELQNKPPPSDDLT
		1		1		TLPSPLILTATVDNANILLQNDN
1		1		1		AHLAAADFRTKFETEQALCLSV
1		1				EADINGPCRVLDELTLARADPE
						MHIENLKEELAYLKKNHEEMN
	1	l				APRGQVGGEINVEMGAAPGVD
		l			i	LSRILNEMREOYEKMAEKNRK
		l				DAEDWFFSKTEELNREVATNSE
		i				LVOSGKSEISELRCTMOALEIEL
ļ						QSQLSMKASLEGNLAETENRY
İ		l				CMQLSQIQGLIGSVEERLAQLL
		l			ł	CEMEQQNQEYKILLDMKMRLE
		1				LEITTYHRLLEGEDAQTVQHTP
						GSPCVPEWAHDRCSLPAOLDE
						WSCOAORGICGHRRENCWERIF
						SOSCFYRPTARSLTSLLGTMST
						CSHQFTSSRSMKGSCSIGGSIGG
						GSSRISSVLAGGSCHAPSTYGG
1						ASCGLGGGYGGGFGSSSSFASG
		ŀ				FGGGYGGGLGAGFGGGLGAGL
	1	1				GGGFAGGDGLLVGSEKVTMOH
		1	1			LGDRLASYLDKVRALEEANAD
						LEVKIRDWYQRQRPSEIKDYSP
					}	YFKTIEDLRHKIIAATIENAQPIL
		l	İ			OIDNARLAADDFRTNYEHELAI
	1					OOTVEAGVNGLCREMLALRGO
23676	54044	A	23811	222	548	TLPCGDHHHSODPARLPSWHD
25070	5-10-1-1	l.,	23011		5.0	LLOERSEKCGSQDWQRDSVQP
				1		GVSYOFCCVPWPAGRSPOPRH
	1		İ			CWGCCGTEASYDSGI*SLSTPG
1		1		1		GGQLQGSV*GPDGAGLQNSHR
23677	54045	A	23812	2	165	SSQLQSS. S. BONGBQ. ISSUE
23678	54046	A	23813	I -	665	
23679	54047	В	23814	ti –	1673	
23079	D404/	lo	23014	11	10/3	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence		09/540,217	coden for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23680	54048	Α	23815	1	331	ARQLRTSAMTMPVNGAHKDA
		1				DLWSSHDKMLAQPLKDSDVEV
						YNIIKKESNRQRVGLELIASENF
		l				ASRAVLEALGSCLNNKYSEGYP
		ı				GQRYYGGTEFIDELETLCQKRA
		l				LQAYKLDPQCWGVNVQPYSGS
		l				PANFAVYTALVEPHGRIMGLDL
						PDGGHLTHGFMTDKKKISATSI
		l				FFESMPYKVNPDTGYINYDQLE
		l				ENARLFHPKLIIAGTSCYSRNLE
						YARLRKIADENGAYLMADMA
		l				HISGLVAAGVVPSPFEHCHVVT
		1				TTTHKTLRGCRAGMIFYRKGV
		1			}	KSVDPKTGKEILYNLESLINSAV
		1				FPGLQGGPHNHAIAGVAVALK
		1				QAMTLEFKVYQHQVVANCRA
		1				LSEALTELGYKIVTGGS/CVISG
l		1				GAGKPG*FSHQDGTGTSQDGG
						RRPFLHHTLSY
23681	54049	A	23816	3	1709	SHPTPRETPTGVEGQATPSVLG
						AAPASQAKATPTLGRPHASDR
		1				GEVAQLDEWSCQAQRGICGHR
						RENCWERIPSQSCFYRPTARSLT
		l				SLLGTMSTCSHQFTSSRSMKGS
						CSIGGSIGGSSRISSVLAGGSC
		1				HAPSTY/GGGLCVSFSRFSFGGA
		l				CGLGGGYGGGFGSSTNFWS\GF
		l				GGGYGAGLGAGFGGGLGAGL
						GGGFAGGDGLLVGSEKVTMQH
		1				LGDRLASYLDKVRALEEANAD LEVKIRDWYORORPSEIKDYSP
						, ,
		l			,	YFKTIEDLRHKIIAATIENAQPIL OIDNARLAADDFRTKYE\HELA
		l				LRQTVEADVNGLRRVLDELTL
		l				ARTOLEMOIEGLKEELAYLRKN
1		l				HEEEMLALRGQTGGDVNVEM
		1				DAAPGVDLSRILNEMRDOYEO
		l	i			MAEKNRRDAETWFLSKTEELN
		l				KEVASNSELVOSSRSEVTELRR
1		l	ĺ			VLQGLEIELQSQLSMKASLENQ
1		1				L\EETKGRYCMQLSQIQGLIGSV
1		l				EE\QLAQLRCEMEQQSQEYQIL
						LDVKTRLEOKYCHPTAALLEG
1		l				EDAHLSSQQASGQSYSSREVFT\
		1				SSSSSSRQTRPILKEQSSSSFSQ
23682	54050	A	23817	2	365	0000000KQTIGTEREQ0000F0Q
23082	D4030	14	2301/	<u> </u>	303	

	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23683	54051	Α	23818	1	464	RRWRICSTWTRSFATAWLPPAR
		1				SANFPRTDRSQGDRGAPAGFAL
		l				APILEFFLWDTSSQKFLQKPHCF
		l				MHGRERAKMGRRAQQESAQA
		l				ENHLNGKNSSLTLTGETSSAKL
		1				PRCR/QGGWAGDSVKASKFRR
		l				KASEEIEDFRLRPOSLNGSDYG
						GDIP
23684	54052	Α	23819	384	805	
23685	54053	Α	23820	3	392	MKYSAIQTLDGEIDLKLLTKVL
		ı		1		APEHEVR*VQWWQQFPGLSTL
						AIPINPAELPGLALCLP*RPWNG
						REKVSLQLREKALSLSYPAKPF
		1				LIEDDVGWDWDHLFTEVSSEV
						LTE\WDPLQTEKEDPAGQARHT
23686	54054	A	23821	63	785	GPIRGPWGPGFGGWGGATPPAF
		1				TSLRRLRAAGLAPRRRAKMGR
		1				RAQQESAQAENHLNGKNSSLT
		l				LTGETSSAKLPRCRQGGWAGD
	l	1				SVKASKFRRKASEEIE\DFR\LRP
		ł				OSLNGS\DYGGDIPI\IPDLEEVQ
						EEDFVLQV\AAPPSI\QIKRVMT
						YRDL\DNDLM\KYSAIQTLDGE
						D/IDLKLLTQK/VLAPEHEVRER
		l			ł	NPSWQDDVGLGTWGPICSTEV
		l			1	SLRRSFTEWDP\LQTEKEDPGG
					1	Q\ARHT
23687	54055	Α	23822	1	441	MSFNCSTRNCSSRPIGGRCIVPV
		1				AQVTTTSTTDADCLGGICLPSSF
		l		l		QTGSWLLDHCQETC/CGCQPLG
		l				GISSVCQPVGGISTVCQPVGGV
		1				STVCQPACGVSRTYQQSCVSSC
						RR\PAKCVGASERIKTPRPASCF
		1				QDLPACCLSLNSSS
23688	54056	Α	23823	1	538	PTRPLQQATCSPLSDIMSFNCST
						RNCSSRPIGGRCIVPVAQVTTTS
			l	1	1	TTDADCLGGICLPSSFQTGSWL
		l	l	1	l	LDHCQETCCEPTACQPTCYRRT
			l	1		SCVSNPCOVTCSROTTCISNPCS
	1	1	l	1		T\PTAGRSP/CVSSGSQPLGGISS
		1				VCOPVGGISTVCOPVGGVSTVC
	[l	l		QPACGVSRTYQQSCVSSCRRTC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:		Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23689	54057	Α	23824	832	1885	AEWDAADKAHVGPGRSPRQPL
						PHVGRRRAAHTTGRGRRTAGC
		1			İ	APAPARTLCILSILGPWCVLPHL
		1				PTLCLGQAQAASCSQGAHGCF
		ı				AGCSCT/GTSASHPV/CCKCPLH
		l				PVCTLADVHGCITSGFSVISLPQ
		l		l		TCAPWTSWRKWSLSDSWQVD
		l				ACPESCCEPPCCATSCCAPAPCL
		1				TLVCTPVSCVSSPCCQAACEPSF
1		1			i	COSGCTSSCTPSCCOOSSCOPA
		l				CCTSSPCQQACCVPVCCKPVCC
		1	İ			VPVCCKPVCCKPICCVPVCSGA
l		1				SSSCCQQSSRQPACCTTSCCRPS
						SSVSLLCRPVCRSTCCVPIPSCC
1		1				APASTCQPSCCRPASCVSLLCRF
1				ĺ		TCSRLSSACCGLSSGQKSSC
23690	54058	A	23825		3097	MDEKRTRLOPEEADVPAYPKV
23090	54056	<u> </u> ^	23023	ľ	5077	ATVPQSEKKPSKYPLCSADAAV
		1				STTPPGSPPPPPPPPPPPVLKVLS
		1				SLKPAAPSPATAATTAAAASTA
l		1				ASSTASSASKTASPLEHILQTLF
1						GKKKSFDPSAREPPGSTAGLPQ
						EPKTTAEDGVPAPPLLDPIVQQF
		1				
		1				GQFSKDKALEEEEDDRPYDPEE
	1	l				EYDPERAFDTQLVERGRRHEVE RAPEAAAAEREEVAYDPEDETI
		l				
		l				LEEAKVTVDDLPNRMCADVRR
23691	54059	A	23826	67	442	NSVERPAEPVA VKFFCLYLQSNEKPMKNFKQG
23091	34039	I ^A	23820	0/	442	SNNQLLMRSFWLLCGIDYKKPP
		l				
1		l				PPPAPGPESVAPALSGPPPPAHS RGPRAVVRA\PPPPGPRHRSPAE
1		l				
		l		i		PPPPPGTEPSVATYSIKASKRVS
22.602	54060	١.	02007		660	QQDRCHNLMKSQE
23692	54060	A	23827	3	657	GERSRRWGPACGPGAAEKTPP
1		l				CDAGAAGAAPGAPGHAGNPGL
İ		l		İ		GSGPDYD\GEATCMHLHPDDPC
Ì		l				PAGAGCCPGSFEVAFPDAAEK
1		l				MKKSLHSLKKGCYSRVCDLPL
		l				DCPGCDSDSGRQAMFSCIVNFQ
ŀ		l				LPKEESPIPGSSQEESSGPSHSEC
		l				DSVGVDWRGQRVSRKLQAPNH
1		1	1	I		SLVLLVTSLFQETTKSHLNRTN
		L	L			DAVSPRKFKRLGALYWGKVEA
23693	54061	Α	23828	141	246	
23694	54062	A	23829	1	609	
23695	54063	Α_	23830	1619	1752	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first eodon for peptide sequence	codon for last amino acid of peptide sequence	*-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
23696	54064	Α	23831	54	470	CSTMNPSEMQRIAPPRRQRHRS RAPSAHKMNRMVMSEEQMKL PSTKKAEPPTWAQLKLTQLA KKKLENTKVTQTPENMLLAAL KTVSTVSAGVPSSSEESDHRER AMMTTVVLSKRRGKCGEKKEI SDCYCVYVERS
23697	54065	A	23832	213	572	IQWYPYQDTIDFLHRIRKNFFKF HMEPKKSLHCQDNPKPKEQSW MHHTT*LQTMLQGYSNQKRMP PHLGSEERLCPAAPSGR*GAPLP GRCAIFQV*SDSLSAGVPNSSEE TATIENGP
23698	54066	Α	23833	296	583	
23699	54067	Α	23834	2452	4557	
23700	54068	Α	23835	3	266	
23701	54069	А	23836	31	499	ARVTAAVSAAAVAKRVWRDG FDCSGYYSLTTILLVQPTKRPIE GRTYADYESVNECMEGVCKM YEEHLK\RMNPINGPSI/THNTTS QLFDFIDVDLADLSCLVYRAUDT QTY\QPYN\RDWIKDEDPTVLLS SGKAQQAREIIVVGSHWGGWG GAWDTGV
23702	54070	Α	23837	3	323	
23703	54071	Α	23838	366	892	PCVYSOFPAGEQCIKLHSPAEA SPPALEASEAQTRKAPECGSGV PEVAGGSCPCLLLCLPRHQAVP TQGFGTYSPCTSHQPHIFPRPAA PHHLGLLQNPINAASCIQCLYP AGVATTMPRRKAEGDAEGD/K AKWKDEPQRRPAKLSAKPAPPN PEAKPKNAPGVTLSIKGTATRF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nuclcotide	Nuclcotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23704	54072	A	23839	201	1412	LWKSRLTFKLAMSRVPSPPPPA
23/04	34072	^	23039	201	1412	EMSSGP\VAESWCYTOIKVVKF\
						SYMWDPSITLAFCREEMG\EV\I
				l		KSSTFSSGANDKLKWC\LRVNP
						KG\LDEESKDYLVTFTCYLVSC\
						PKSEVRANFKFSILNAKGEETK
						AMESQR\AYRFVQGKDWGFKK
						FIRRDFLLDEANGL\LP\DDKLTL
						FCEVSVVQDSVNISGQNTMNM
		l				VKVPECRLADELRGLWENS\RF
						TDCCLCVAGHEFQAHKAILAA
						RSPVFSAMFEHEMEESKKNPSL
			ŀ			KSMDV\EPEVFKE\MMCFIYTG
						KAPNLTKMA\NDLLAAADKYA
					l	L\ERLKVMCEDALCSNPV/SVEN
						AAEILIL\ADLH\SADQLKTQAV
						DFINYHA\GDVLETSGWKSMVL
						SHP/HNLVAEAYRSLAS\KQCPF
						LGTPTQNALKQSLKILLCC
23705	54073	Λ	23840	I	575	FRHRKIIY*NLLKQEDSINVTEM
						TSIRASVDILIPDPPPKDDEMDT
				l		DKQEKKEVPKCGFLPGNEKVL
						SLLAMVKPEVWTLKEKCIVVIT
		1				WIQHLIPKIEDGNDFGVAIQEK
		1			ļ	VLERVNAVKTKVEAFQTTIS/K
						MDYRALVHERDEAAYGELRA
						MVLDLRAFYAELYHIISSNLEKI
						VNPKGEEKPLMYPTRD
23706	54074	A	23841	1	1092	
23707	54075	Α	23842	130	397	EGVHSQVNLFPIKGNQTHVPAS
						HGTDSALPPTLQMDYRALVHE
						PD*AAYGELRAMVLDLRAFYA
						ELYHIISSNLEKIVNPKGEEKPF
23708	54076	Α	23843	20	334	VRLPGSTHAGGDRSSMAKPCG
					l	VRLSGEARIOVEVLRONLFOEA
					l	EEFLYRFLPOKIIYLNOLLOEDS
						LNVADLTSLRAPLDMPIPDPPP
					l	KDDEMETDEOEKKEGK

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last eodon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
23709	54077	A	23844		898	MKQKVAAEAGRGGII.QAKVLP SGTLSRLAEALFGVQTYDQDA YGFLTPSAQSMKVFCCLWKLV CKQLEGDVAQGLAIDGDVKKH SGLEWQVETIQNLVDSYMAIV NKNMQDLMPKTIHLMINNTKE FIFSELLANLYSRGEQNTLMGE SAEQAQGREMBREMHHVLRE APSIIGDIKTTIVSTPTGSPKSSPT TQCRLPAVPPARPGLRSPAPGPL PAASTLGEAPVVPSTPGASPDPL GPPPQVPSRPSRGPQPQDRMPC GLAGDLTYGFLEPTLQAIIRWS LLLRPPANSFEQS
23710	54078	A	23845	99	429	FSDAAAVARQPSFGKALGAPQP ADTQPTPSMPPRCPGGSETKPK KAAGKDKSSDK\KNKTKKQMK AEKGGKEKQAEVADQETKDLS AENGDTENKESPASDEATEKDT KSD
23711	54079	A	23846		390	TWMFIAAVFRIASTWKQPRYFS VGKQDQKRTSRKLQQTCSRGA *WLEGKLTNRKE*HQYQRKGH PRKNTIQRSPTLKTKANIILNGQ KLEAFPLKTGTRQRCPLSPLLFN MVLEVLARSIRQEKEIKVFK
23712	54080	A	23847	522	733	ASAADTQANRVWSGPP/RKLQ QTCS*GS*LLQGKLTNRKDIHT KTPSVHHHHQRPKTLNQQRSK ETKKAIT
23713	54081	Α	23848	46	330	
23714	54082	Α	23849		873	MAEKKDPTKEPEKKHLLA WM NKLLVELDPKYNDSNTDGSA VYHKIIIACKPVSPPPSAMIGSF PRPSSEADAGAMI.CVQPARTKI LGILGLKATSLQYLRLHTGKPL LPKIDESTKMGRNQPKKAENSK NQNASSPPKDHNSSPVREQNR MQNEFDELTEVGFRRDRSSK NSVVIKALPRHLINGGRSQEWR LGIGARDSFKTSGRRGKSPAGP KTTYPEDEQA/GSRPRPIPTRQ ADPRTPPPTRSEKE/AQLSLATD QMCPSLAPLIK*PQES**WKQRI SI*MEHL
23715	54083	A	23850	1075	1350	AIITANVFKFVSIDRGLVCNMEL EGSNCFIDVCSWTSEMKSKGTS RR*KIT*GNSSGSLPRDTQRSGH RNSSLCCERSKYGGSRVDIRRK VF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
23716	54084	A	23851	1	956	MEVLARVFRQEKEMKGIHIGK
		1			1	EEHKLFLFADNMTLCLEKTKDS
						VKKLVEQPSSFGSVDWLCLSSC
						SGSTHTPRPADVSPGSLPGPGQ
					Į.	TFGTREPPQAVSTKEASSSNLH
				i		APERTVAGLTFTTEQVRALEGV
						FRHHQYLGPLERNWLAREMQL
	ł	ı				SEVQIKTWFQNRRMKHKRQM
						QDSQLNGPLSGSLHGPPAFHSP
		1		ŀ		SSGLANGLQLLCPWAPLPGS/L
		1				WLPLGSFWGLRQMDQEALASS
						WASCCGQPLAYHPPRPGSGAH
		1		1		KLGPALSTGPRGLCALPETGDA
1		1		1		FEETPPTTCLPHKRSVQIASDAY
	ŀ	1		1		PEDSAVLFTSWRHLSP
23717	54085	A	23852	1	846	MEAFDANYTFQSGPSAAGLLV
		1				FAGGPLQALCAWVSAVEAAEQ
	İ					WILQPSSFGSVDWLCLSSCSGST
		1				HTPRPADVSPGSLPGPGQTFGT
						REPPQAVSTKEASSSNLHAPER
				1		TVAGLTFTTEQVRALEGVFRHH
	1	1		1		QYLGPLERNWLAREMQLSEVQ
	1	1			İ	IKTWFQNRRMKHKRQMQDSQ
				ŀ		LNGPLMGSFPGAPWLSTHHFLA
				1		LAKGLQ\LLCPWA\PLPGSPGCP
		i		1	i	LGSFWGTP\QMDQEALASSWA
					İ	GCCG\QPLAYHPPRPGSGIDLPK
						GTQLSLRLFQDISSTRGDHRSG
23718	54086	Α	23853	3	239	
23719	54087	Α	23854	10	1154	RRAGFGGVRASGAMGTPPGLQ
ļ				1		TDCEALLSRFQETDSVRFEDFT
						ELWRNMKFGTIFCGRMRNLEK
						NMFTKEALALAWRYFLPPYTF
				I		QIRVGALYLLYGLYNTQLCQPK
	1			1		QKIRVALKDWDEVLKFQQDLV
	1				1	NAQHFDAAYIFRKLRLDRAFHF
				l	1	TAMPKLLSYRMKKKIHRAEVT
ļ						EEFKDPSDRVMKLITSDVLEEM
				1		LNVHDHYQNMKHVISVDKSKP
	1	1		1	1	DKALSLIKDDFFDNIKNIVLEHQ
1	1	1		1	1	QWHKDRKNPSLKSKTNDGEEK
1		1			1	MEGNSQETERCERAESLAKIKS
	1	1			1	KA\FSVVIQASKSRRHRQVKLD
1	1					SSPLDSASGQGQVKATRKKEK
	1			1		KERLKPAGRKMSLRNKGNVQN
						IHKEDKPLSLSMPVITEEEENES
1	1	1	1	1		LSGTEFTASKKRRQH

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23720	54088	A	23855	16	447	INNCDKANCSTTCFNGGTCFYP GKCICPPGLEGEQCEISKCPQPC RNGGKCIGKSKSVCEPGCGAH GTCHEPNKCQCQEGWHGRHCN KRYEASLIHALRPAGAQLRQH TPSLKKAEERGDPPESNIYIWVE LPTSGNVLSYTKFI
23721	54089	В	23856	103	1989	
23722	54090	A	23857	3	1295	SGRGYSRHMADYLISGGTGYY PEGGITAQQLFTGAAARDRLT YNNFLILPGFIDFIADKYDLTSA LTWKITLKMPLISSPMDAVTEA DMALAMALMGGIGFIHHHETTE FQANEVRKYKKFERGFIMDPV VLSPSHTYGDVLEGKMRIGFS GUPITETGTMGSKLDSHKQLLC GAAVGTREDDKYRLDLLTQVG OVDVIVLDLSQGNSLYQMAMYY YIKQKYPHLQVIGGNWTTAQ AKNLIDAGVGLHYGMGCSSIC ITQEVMAYSWPQGTAVYKVAE YARHFDHGEHGKALAGASTV MMGSLLAATTEVPSKYFFSDG VWLKKYRGMGSLDALEKSSGS KQYFMEGDKLKIAQGYSSIQ DKGSIQKFVSYLLAGIQHGCQD TOYCNLSVLLSMMYSGERKFE KWTMLAQIKGGVQGLHSYKK
23723	54091	A	23858	770	1008	LAPSPCVRSKGRKESPDSR/RPL WLRPIAPALGPSGRRGLAPDSR LNPHRSLLGTGNYDVNVIMAA LQGLGLAAVWWDRRR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nueleotide insertion)
				sequence		
23724	54092	Α	23859	I	1489	MTRWSSYLLGWTTFLLYSYESS
						GGMHEECVFPFTYKGSVYFTCT
						HIHSLSPWCATRAVYNGQWKY
						CQSEDYPRCIFPFIYRGKAYNSC
		l				ISQGSFLGSLWCSVTSVFDEKQ
		l				QWKFCETNEYGGNSLRKPCIFP
		1				SIYRNNVVSDCMEDESNKLWC
		l				PTTENMDKDGKWSFCADTSNL
		l			1	GMGVGPTPRHHRRASTRHSRPP
					1	PADPDGGGERPRTADDRQDDA
l		l				TEDDGTRCRGRQTRTQATNAN
l		l				ANHKKSNRTKPPPRHKPNKAK
						RHTKKETPQNT/CGTQDKNPDS
						GRPKLFRRACITSTGTRGELWT
		1				RSREHPAK\QGRRSGKQRLRPW
l		l				DRMLSPDWHRAQRIAQVWKPE
l						QEDGLSIGFGDQPGPHSDTSSR
		ŀ		İ		QQVRRSKATEMHVGGRVDFD
		ŀ				DFVELMTPKLLAETAGMIGVQ
						EMRDAFKEFDTNGDGEITLVEL
						QQAMQRLLGERLTPREISEVVR
						EADVNGDGTVDFEGDIDGAWG
		1				RHGTQGPHWPWGLGDVRGEPL
23725	54093	Α	23860	283	490	GDIWMGTQPKPYHS\PPGPSPN\
		1				PCPPHFKNQSMPSPKSPPKVFNS
		l				FPALTQKVHTSKVSSETRQVPS
						HL
23726	54094	С	23861	82	222	
23727	54095	Α	23862	27	454	VDGRIAGEITRRGSRARPRPGPQ
						CPPGRPGTAMIKAILIFNNHGKP
					1	RLSKFYQPYSEDTQQQIIRETFH
						LVSKRDENVCNFLEGG/LVFVE
					1	TLDKCFENVCELDLIFHVDKVH
						NILAEMVMGGMVLETNMNEIV
						TQIDAQNKL
23728	54096	Α	23863	1	403	RGSRARPRPGPQCPPGRPGTAM
		1				IKAILIFNNHGKPRLSKFYQPYS
		ŀ				EDTQQQIFRETF/HLGSKKEEKG
]						WNFLKGGLLIGGSDNKLDYRH
		1			1	YATVYFVF\GGDFSKRELGFLN
		_				LIQEFGEPLDKCFEKVWELDLIF
23729	54097	Α	23864	215	599	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23730	54098	Α	23865	2	683	GRIAGEITRRGSRARPRGPOCP PGRPGTAMIKAILIFNNHGKPRL SKRYDPYSEDTQQIIRETTHLV SKRDENVCNFLEGGLLIGGSDN KLIYRRYATULYPVECVDSSES ELGILDLQVFVETLDKCFGNV CEL\DLIFHVJDK VHNILAEMV MGGMVLETMNIEIVTPIDAQN KLEKSEAGLAGAPAIRAIVSAV KKNML/REIPRINIGIDSIKIVP
23731	54099	В	23866	94	726	
23732	54100	В	23867	94	I I 46	
23733	54101	В	23868	1	360	
23734	54102	В	23869	1	1371	
23735	54103	В	23870	1	639	
23736	54104	В	23871	280	2391	
23737	54105	Α	23872	3	244	
23738	54106	В	23873	103	285	
23739	54107	A	23874	70	378	MPLRILLMHGEEEEEEEEEE EEEEEEEEEEEEEEEEEE EEEEEEEE
23740	54108	A	23875	5	156	RKKGGG/EREEEGEGEEEEEE EEEEEEEEEEEEEEEEEVVL LTPGTGAEDFKKH

SEQ ID	SEQ ID NO:					Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23741	54109	Α	23876	1	1808	MEVLSIDNLAGHKVFYQKESHS
			l			WKQQKQDLMSGKRLSAANPLF
		1				ALLLPVLWTTGQAPSPGPGGDP
						ILOLWDDAOOMAGSPRTKKKG
		1				CTRNSQAARLSFOSRGGOGSLT
		l				EKMIFEORFGGGKGESRKDAIA
		1			1	IIVVDIVTLQPCRTVEWVPTRPF
		l				RSLQGHFQGGLRRQQMMVME
						GVWVLSCAMDLTSPRDCLPFC
						VSLPVLLLSAPLCSCSTAGGSPV
i					1	TSPHHFPVEGGLGRVRRCYINE
						PVVTTLGWESVCWVSFWLLLR
		l		1		IRYSTSGQAACHAESQELNPRG
ĺ	ļ.	1		1		FGCLGHRNVTWVKCLLPALQP
		ł				EGSVPFLELLEPFIFASASKTFV
						AANLVTWVACFLLVQAAFQLS
		1		1	Į.	EYRQGVGIPVADRTGDAPTLGL
		1				SYEADVAEGHGFFLDQESVIQII
		1				PEPLPOFYLCLLDAELTGVOGE
						PALPSMQIFSSPEAATGVHNDR
		1				GAEHFPVSRQPSAELGLRPMQA
		1				SOERKINFPGPSPGFPPKRONKT
l		l			1	LSRREKKEEEKEEKEEEEEE/E
l		i				GEEEKEKEEEEEKEEEEEEEE
ļ		1				GGEEEEEEEEEEEEEEE
1		1				KWLYOYDPEDKAQSKOWLSR
		1				GGSGPVKAKADWWRAKEIHSL
		l				OPROVIPGIPSM
23742	54110	A	23877	459	636	IPGGSNGLEAEGPPPPPLPVAAV
		1.	2007	1.07		LA/AGSGGLAVASCAFTICALR
	1					WPGVPCSRVPIWTPS
23743	54111	С	23878	183	254	
23744	54112	A	23879	103	335	LGSGDLPWEINPLSSCSLLHEK
						DPPTTSGPQTDRPKEHLINFKSG
l		1				ACYTCRKSGH*AKECLQPGIPIK
						PHLICVGPH
23745	54113	Α	23880	159	306	LGSGNLP*EINPLSSCSLFREEDP
		ł				PTTSGPQTNQPKEHLTNFKSAA
		l				ED
23746	54114	С	23881	80	106	
23747	54115	Α	23882	317	427	
23748	54116	A	23883	353	511	
23749	54117	В	23884	I	320	
23750	54118	A	23885	1	624	
23751	54119	Α	23886	224	355	
23752	54120	A	23887	872	1003	
23753	54121	Α	23888	I	603	
23754	54122	Α	23889	2	214	
23755	54123	Α	23890	459	601	DVDRHVRGSNFHHNEIRSLAAT
	l	1		i		PRAPGTLAQ/GLTDSFPDLLGLA
		L_		L	l	AED

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	deletion, \=possible nucleotide insertion)
23756	54124	Α	23891	2	267	QWDGWGARKGMEWEDDLPLE FGRPTASLLSDRPQPNSSRGSDV PFL\PPSLPCHSSACLAATPRAPG TLAQGSLTDSFPDLLGLAAED
23757	54125	Α	23892	72	299	
23758	54126	В	23893	1	1470	
23759	54127	A	23894	134	1276	
23760	54128	A	23895	2	847	PPPPRVLRTGGGASLTAQAPGT PRRPQEGEPVTPGRSLGLRSSCQ ASVSLLLPVIVACASLEPGRGR AFRDEGAWTTRVSAAVISSQA ALLSSILWSRRLSFSAHTHAFG TVALGEYAFRMFIRLSARLFGIE VTRPTNS/RSLMKVVKLFSEJW PLAKKETYDWYPPHIITY/AL ELMAGRLRFLGTLTEDEAFRDF MGMEAKTDLKEAFVGKGGNP KEREKGKRAAKKGNSCLVPSR RGDPLSSVAERRKLHLLSFPHIG GDVIFLMGSLFGRYTSHLGEI
23761	54129	Α	23898	3	1224	MRRNKPRORGVREARPAGGAG PFWRGARCSGAGEGGGERRG SPPAALALAPASGPRRNFPPDA RCLIYQQIHQGAFLLAGSASLS AVTGYPEGGARTEDVCPEK MFGFHKPKMYRSIEGCICISGA KSSSSRTDSKRYEKIDPQSCFG LHETRSGDICNACVLLVKRWK KLPAGSKKINWNHYDARAGP SULKTTFEPKIKVKTLSGINRIKS TQISKLQKEFKIRSNSDAHSTTS SASPAQSPCYSNQSDDGDTEM AGSSNRTPVSFLDLTYWKRQK ICCGIIYKGRFGEVLIDTHLFKP CCSNKKAAAEKPGQPEPLPIS TQEWTECPDLSLVVSWRPLS LPAGCSLHGLAADWTTDMITV TTSKCREPSCQQEGSFQRKAF PKRLPKMAEVQVLVL
23762	54130	A	23900	69	355	
23763	54131	А	23901	316	830	VSLSQTLHPPMTLQLSRYSRGW GLSSDRPRGAKAKVPSCFGTAP RIGPEAAPRLLAHHLVPAGRGS/ ARAPAPGWRLGSP\GAGRKGR DFRLPASGLAVSGEWDPIPWGY VGEVGGCSCQWFSPGVQQKRY RRKVWLGQPQDLPGPSALADK GPALAKADGQRVNVASVGLQP

de hod				
.	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop eodon, /=possible nucleotide
١	09/540,217	sequence	or peptide sequence	deletion, \=possible nucleotide insertion)
A	23902	I	1044	MSYKVTQEISKELRWISAVLVT
ı				WDLRGETGTRVCIAMGVSRMQ
				YSIMAKIVGSGNNLELNPAADK
ł				LDNLGAQCGLASVKVSARAAP
- 1				PFAPATLVALLEARAAELPLPPP
				PGVSLRCVEAPDVRGRGRGGG
- 1				EGSGVTRLLLPPPPPSGASFLQR
- 1	1			GRPVTPALFQAGWGRAADRPG
- 1				PADLAGGAGGRSGGRRMLPRS
				PGAALRDRSALEDGIVEGYGPL
				GEPEEGSRSSFLESKVAFNYCN
ŀ	ļ			KVAPLDFGNEAVEQCHTMSDR
- 1				KAVIKNADMSEDMQQDAVDC
- 1				A\TQAMEKYNIEKDIAAYIKKEF
				DKKYNPTWHCIVGRNF\GSY\V
				PHTEDKHFIYFYLGQVAIL\LFK
A	23903	33	427	
Α	23904	72	777	EWKQAGEGGGGQRRGSPPAVL
				ALAPASCPRREMSYLPTNPPGSI
				LICWLGVSLSSHRSAGGGSAIY
				G\SYYPEEKMFVFHKPKMYRSI
				EGYCICRAKSSSSRFTDSKRYEK
- 1				DFQSCFGLHETRSGDICNACVL
1				LVKRWKKLPAGSKKKTGIMCN
				OSDDGSDTEMASGSNRTPVFSF
				LDLTYWKRQKICCGINYKGRF
				GEVLIDTHPFKPCCSNKKAAAE
				KPEEOGPEPLPISTOEW
Α	23905	234	989	PKTSDOAYYCPEEKMFGFHKP
1				KMY\RSIEGC\CISGAKSSSRFT
				DSKRYEK\DFQ\SCFGLHETR\SG
				DI\CNACVLLVKRWKKLPAGSK
				K\NWNHVVDARAGPSLKTTLK
l				PKKVKTL\SGNRIK\STQISKLQK
				/ENFKR\HNS\DAHSTHPOVPSP\
	1			AOSPLFTVNOFRWTGSDTGVG
	1			FPGSNRNHPVFSFLDL\TYWKR
	1			OKICCGNYKGRFGEVLIDTHLF
- 1				KPCCSNKKA\AAEKPEEQGPEP
1 _				LPISTOEWVTEVFM
A	23906	3	1059	
	A	A 23903 A 23904 A 23905	A 23903 33 A 23904 72	A 23902 1 1044 A 23903 33 427 A 23904 72 777 A 23905 234 989

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23769	54137	A	23907	1	1697	MRPCALCWRSAVTAAPERAVL
		l				ESRGGAPDTPRRLHSFTQTRSLS
						PQAHAGPHTRLHPRQGHAHMR
						GGGRVTDTRRHIPNADVPRYK
						LTCTGGVTTFVALYDYESRTET
		1				DLSFKKGER\LQIVNNTKGDW
						WLAHSLSTGQTGYIPSNYVAPS
						DSIQAEEWYFGKITRRESERLLL
			ŀ			NAENPRGTFLVRESETTKGAYC
		i				LSVSDFDNAKGLNVKHYKIRK
						LDSGGFYITSRTQFNSLQQLVA
İ						YYSMSHCPPFSAAGWYGGGCP
						EEPQKHADGLCHRLTTVCPTSK
						PQTQGLAKDAWEIPRESLRLEV
						KLGQGCFGEVWMGTWNGTTR
						VAIKTLKPGTMSPEAFLQEAQV
		l				MKKLRHEKLVQLYAVVSEEPI
	ľ					YIVTEYMSKGSLLDFLKGETGK
		l				YLRLPQLVDMAAQIASGMAYV
						ERMNYVHRDLRAANILVGENL
						VCKVADFGLARLIEDNEYTAR
İ						QGAKFPIKWTAPEAALYGRFTI
						KSDVWSFGILLTELTTKGRVPY
	ŀ					PGMVNREVLDQVERGYRMPCP
						PECPESLHDLMCQCWRKEPEER
						PTFEYLQAFLEDYFTSTEPQYQP
23770	54138	Α	23908	1	736	MDPGAALQRRAGGGGGLGAG
						SPALSGGQGRRRKQPPRPADFK
		1				LQVIIIGSRGVGKTSLMERFTDD
						TFCEACKSTVGV\DFKIKTVELR
						GKKIRLQIWDTAGQERFSSITSA
						YYRSAKGMILVYDITKKETFDD
						LPKWMKMIDKYA\SEDADLLL
			l			VGNKLDCETDREITRQQGEKFA
						QQITGMRFCEASAKDNFNVDEI
		ĺ				FLKLVDDILKKMPLDILRNELS
						NSILSLQPEPEIPPELPPPRPHVR
23771	54139	В	23909	84	441	
23772	54140	A	23910	6	254	
23773	54141	A	23911	1	230	
23774	54142	A	23912	3	187	
23775	54143	A	23913	3	143	
23776	54144	A	23914	3	481	
23777	54145	Α	23915	1	711	
23778	54146	Α	23916	55	381	

SEQ ID	leen in vo.	Alot	SEQ ID NO:	Nucleotide	Nucleatide losation of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1.0.	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		L				
23779	54147	Α	23917	240	875	GLSAESTPTSIMSMTLGYWDIR
						GLAHAIRLLLEYTDSSYEEKKY
	i					TMGDAPDYDRSQWL\NEKFKL
	i					GL\DFPNLPY\LIDGAHKMHPEA
						TPFLCYI\ARKHNLLLGETGRRE
İ						KISSWDIL\ENQAMDVSNQLAR
		1				VCYSPDFEKLKPEYLEELPTMM
						OHFSOFLGKRPWFVGDKITFVD
1		1				FLAYDVLDLHRIFEPNCLDAFP
						NLKDFISRFEVSCGIM
23780	54148	A	23918	64	774	CLSAESAPTSTMPMTLGYWDIR
25700		Ι'	23310		l'''	GLAHAIRLLLEYTD\SSYEEKKY
1					l	TMGDAP\DYDRSQWLNEKFKL
1						GL\DFPNLPYLIDGAHKITOSNA
		1				ILRYI\ARK\HNLCGGDRRREKIS
		1			l	CWDILENQVM\DNHMELVRLC
1		1				YDPDFEKLKPKYLEELPEKLKL
l .		1				YSEFLGKRPCFLG\DKITFVDFL
		1				
				i	ļ.	VY\DVLETK\QVFEPK\CLDAF\Q
	ļ	1				NLKDFISRFEGL\EKISAYMKSS\
22701	54140	١.	22010	162	217	RFL\QELVFSKMAVWGNK
23781	54149	A	23919	163	317	LSPNNYYRKLPQKVKLE*RYHR
						LALFSFPTKKLTPEKVSLCCRFL
		Ļ.		10		EMGYIF
23782	54150	Α	23920	62	255	VRLTAQKSLGPLTSTHSLWGSA
		1				\RARHREPVGSVCGCDPCEQLL
		_				LLVSQLRAPGVDSAAAGRPV
23783	5415I	Α	23921	173	467	KSRFCFFGVVLKLPKKLHIR*WI
		1			\	GATL*NQP*TIFQTDGDCSHQSL
						GPLPACDLWDQLHLRSKQGGS
		1				VCGCDPCEQLLLLVSQLRAPGV
		ــــــــــــــــــــــــــــــــــــــ				DSAAAGRPV
23784	54152	Α	23922	35	548	PRSCPSLSPGLSCLLQ/RLVNYQI
		1				SIKCSNQFKLEVCLLNAENKVV
						DNQAGTQGQ\LPAVGQARWGS
		1				GCRGAHTW\PLAVIGAADCTG
		1				QRGWEGV*AFPGLTEPVSSLFC
		1				GQDGD/CSPHSM*VLSHVVLWL
i		1				LNGLC\SLVLGSVCGCDPCEQL
		1				LLLVSQLRAPGVDSAAAGHPV
23785	54153	Α	23923	I	599	MSPAGLVNYQISVKCSNQFKLE
		1		1		VCLLNAENKVVDNQAGTQGQL
		1				KGRPDGAQAV/VGAHTWRFGC
1	1	1	l	1		RRCG*LHRSHWGL*LLHTPCGR
	İ	1				PHCARHREPVAHSLLCSRAISGS
	1	1			1	EPHFWQCSVQ*PQLCVPGRVAT
1		1	l			PLGELDSGDGDCSHQSLGPLPA
1	1	1	l			CDLWDQLHLRSRQGGSVCGCD
		1	l	1		PCEQLLLLVSQLRAPGVDSAAA
		1		1		GRPV
23786	54154	A	23924	290	528	
25,00	124124	1.	23724	200	520	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23787	54155	A	23925	274	571	KSRFCFFGVVLKLPKKLHIR*WI
						GATL*NQP*TIFQTDGDCSHQSL
				İ		GPLPACDLWDQLHLHSRQGGS
1				1		VCGCDPCEQLLLL\VSQLQAPAI
		1				DSAAAGRPV
23788	54156	В	23926	204	410	
23789	54157	Α	23927	29	330	YLGVELVKSAYPFFQIHLSFPSC
		1		1		MWLCGLFLLFYIDFLTLMCGD
		1				KPLICKKHQLSASTCA/CTIC*G
						ALTLLMEIEQGPCDLSCASREV
						STARSLAELFSGP
23790	54158	Α	23928	259	699	ETSHQVMDRSNPVKPALDYFL
						NRLVNYQISVKCSNQFKLEVCL
				•		LNAENKVVDNQAGTQGQLKV
						LGANLWWPYLMHEHPAYLYS
		1				WEDGDCSHQSLGPLPACDLWD
1		!				QLHLRSRQGGSVCGCDPCEQLL
		ļ.,				LLVSQLRAPG\VDSAAAGHPV
23791	54159	A	23929	245	695	
23792	54160	Α	23930	302	745	ETSHQVMDRSNPVKPALDYFS
		1				NRLVNYQISVKCSNQFKLEVCL
				1		LNAENKVVDNQAGTQGQLKV
		1				LGANLWWPYLMHEHPAYLYS
		1	ì			WEDGDCFTPKALDPLPACDLW
		1				DQL\HLRSRQGGSVCGCDPCEQ
23793	54161	١.	23931	2	293	LLLLVSQLRAPGVDSAAAGRPV VFCHTDLRKNLTVDEGTMKVE
23/93	34161	Λ	23931	ľ	293	VLPALTDNYMYLVIDDETKEA
		1				AIVDPVOPOKVVDAARKHG\V
		1		1		KLTTVLTTHHHWDHAGGNEKL
		i .		i		VKLESADLGIRLF
23794	54162	A	23932	1	441	
23795	54163	Α	23933	1	630	
23796	54164	A	23934	3	265	
23797	54165	Α	23935	3	1062	SAVQPGPVMVVGRGCSRRSLA
		1		1		ALGAACARRGLGPALLGVFCH
1		1		1		TDLRKNLTVDEGTMKVEVLPA
		1	ŀ	I		LTDNYMYLVIDDETKEAAIVDP
1		1				VQPQKVVDAARKHGVKLTTVL
			į.	l .		TTHHHWDHAGGNEKLVKLESG
		1	l			LKVYGGDDRIGALTHKITHLST
			1			LQVG\SLNVKCLATP\CHTSGHI
		1	l			CYFVSKLGG\SEPFVVFTGDTLF
1	1		1			V\AGCGEVLMKGLRDEMCKAL
			1	1		LEV\LGRLPPDTVGSASLRAGR
					1	VYCG\HEYTIKQTSKFAR\HVET
			1	1	1	GNA\AIREK\LPWA\KEKYSIGEP
1	1	1	1	1		TVPST\LAE\EFTYNP\FIKVREER
	1	1	1	1		RCKQH\AGETDPVTTMRPVRRE
		1	1	1		KDQ\FK\MPRDLRPPLHLQQDL
		L.	l			GIKALLDRLF

SEO ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23798	54166	Α	23936	107	750	SHLWSPVSEPLQACSFC/VIRDK
l		i				YRYFACLMRARSEEHTNEKDM
1						MKAIRLLKEAEEEFWYHOHPO
l						PCIFPDSPRGTSYVTHECYKVPE
l						WCLDDWHPSEKAMYPDYFAK
						RKOWKKLRRESWKREVKOLO
İ						EETPPAGKEECKLHEGMNYVV
						LTCIPGAIKQYLACSRYSKTPGP
		l			Ī	RDSCWEVQSDNQCWWWLDSG
		1	1			AHGLVTSFKGVMMTVEATTM
						AAVF
23799	54167	A	23937	3	718	PASTISCSVTRSRRAVSRLSARP
23777	34107	n	23737	ľ	' ' '	GKVSAVMASLASGPYLTHQQK
						VLRLYKRALRHLESWCVORDK
						YRYFACLMRARFEEHKNEODM
						AKATOLLKEAEEE\FWYROHPO
				1		P\YIFPDSPWGHPPYERY\DCYK
						VP\EWCLDDWHPSEK\AMYP\D
						YFAKREQWKKLRRESWEREV\
						KQL\QEET\PPGGSF*LEALPPGP
		1			1	KRKVILPPLWW\YIVTRP\RERP
		1				MLERERPHLSCLQVKYVTEHG
23800	54168	A	23938	269	488	TDGVIFIGLSWSPRRCGLGIORD
23800	34106	^	23936	209	1400	AERIFFALDQASCGFTIHPHIDA
						AETPRGLT\MHLCRRCQRGQSQ
						AEKLP
23801	54169	A	23939	1	1560	MVHLGAPVMALNLOGKHLCD
23601	34109	^	23939	<u>'</u>	1300	STASSLLPRRRESNSDDTARED
						AROKRPANTDMRTOELOTRET
					1	GOGTARETEVATPLSGPSAGDS
ŀ	ĺ					PPAASEMSCGQLVLALLLLFFL
	ŀ					LVWNTGTKSGSGRVARLAQLA
				1		HDCDGGLALGLCYAVLDQRSV
1						VQATPQAPKPGPVQPLTVQGL
1						OPVHVAOESSGNEITARROSAR
		l				SRPRGDRTAPSRRVTRRPKHAP
				•		ERAPIQQGNIVTVRAPTARKKW
		1		1		RPAVGTPTGAHPVCTAKGTGG
1						
		1				RPTPTQAHTRYTERDPIRNSAEL TGSRPCTGRERSGVEACKAAV
		1				
						HNLL\AQTGSSGLIINYPLDD\Q
		1	١,			DAIAVEAACTNVPALFLDVSDQ
	1	1				TPINSIIFSHEDDVWRTPLVAPT
	1	1	1	1		NLNTRSSQPARHHRAGAVRRH
	1	1	1	İ	1	HMTVARSATAHARLRRHAKPQ
		1	1		1	GRERIKPSGAREGSEPLGGESGI
1	1	1	1		1	RTPRGRERDPNPSGAKAGSEPG
1		1	1			RIPETLPRCKGETEAKCWARGN
1						FHTAQGSWPRGGRSRWEPLEA
		_				PALSQQAHGGELPLPAA ARSHR
23802	54170	В	23940	98	229	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	
NO:	nf peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23803	54171	Α	23941	1	256	
23804	54172	Α	23942	289	433	VLWSCAAEDFDWLVYRPIGSA
				l		LSCRLIGPWAIGMWETGFP*VL
				l		WSCAAEDFDWLVYRPIGSALSC
	l					RLIGPWAIGMWETGFPLLSSVS
				1		LS
23805	54173	Α	23943	450	1260	LCACVVGGEKRSGPRIGCAGA
						GKDPFWRAVGVAQKAAAVVA
				1		CGAASPYRNRADVSVSDHSQS
						RGPGGLSLSGGTVCDMGKKTK
	1					RTADSSSSEDEEEYVVEKVLDR
	1	1				RVVKGQVEYLLKWKGFSEEHN
		1				TWEPEKNL\DCPELISEFMKKY
	l	ĺ				KKMKEGGN\NKVREKSESSKR
	l	l	(KSNFSDSAEDIKCKKKREOSND
		l				IARGFERGLEPEKIIGATDSCGD
						LMFLMKWKDTDEADLVLAKE
		l				ANVKCPQIVIAFYEERLTWHAF\
						PEDAENKEKETAKS
23806	54174	Α	23944	3	267	AMVGGGGVGGGLLENANPLIY
						QRSGERPVTAGEEDEQVPDSID
				1	1	AREIFDLIRS*AWPLTPQPAYW
						YPGPSSCQGCYRCFLESLTMRN
23807	54175	Α	23945	1	942	
23808	54176	Α	23946	3	470	AMVGGGGVGGGLLENANPLIY
1		١				QRSGERPVTAGEEDEQVPDSID
l	1		1			AREIFDIRRCWARAGSGGLRW
	1					GEQ*YRAAGGAASQQGVPGRG
	I					F*VGKGAESLFLHFPQ\LIRFLN
	1	l				DP\EHSLTLEELNVVEQVRVQV
		l				SDPESTVAVAFTPTISHWSMGT
23809	54177	Α	23947	2	604	ARSHRISGGGSAMVGGGGVGG
		l				GLLENANPLIYQRFGE\RPVTAG
	1					EEDEQVPDSIDAREVF\DLIRSH
						QMTPEHP\LTLEELNVV\EQVR\
	1				į .	VQVSDP\ESTVAVGFPKPTHSGT
	1					CRHGPPLIGLS\IKVK\LLRSLPS
	1	1			1	AFSRWDVAHLLPGDPLPQEHA
	1	1		1	1	VEQATLQIKEAGWATALGRTH
		1				HLLEVVN\QCLSSPAPGTWAFV
23810	54178	Α	23948	1	1002	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Aminn acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
	<u> </u>	<u> </u>				
23811	54179	A	23949	89	2587	STSRDRRLSKRSRTTMAAQPRG
		1				GDYRGSFLLSILLGTPWEAWAG
						RILYSVSEETDKGSFVGDIAKDL
						GLEPRELAERGVRIISRGRTQLF
İ						ALNQRSGSLVTAGRIDREEICA
		l				QSARCLVNFNILMEDKMNLYPI
		1				DVEHDINDNVPRFLTEEINVKI
		l				MENTAPGVRFPLSEAGDPDVG
1		l				TNSLQSYQLSPNRHFSLAVQSG
		ł				DDETKYPELVLERVLDREEERV
						HHLVLTASDGGDPPRS\$TAHIQ
		1				VTVVDVNDHTPVFSLPQYQVT
				i		VPENVPVGTRLLTVHAIDLDEG
		l				VNGEVTYSFRKITPKLPKMFHL
						NSLTGEISTLEGLDYEETAFYE
1		1		ł		MEVQAQDGPGSLTKAKVLITV
		l				LDVNDNAPEVTMTSLSSSIPED
		ŀ				TPLGTVIALFYLODRDSGKNGE
						VTCTIPENLPFKLEKSIDNYYRL
					1	K\TTKNLDRETLSLYNITLKATD
					i	GGTPPLSRETHIFMQVADTNDN
1					ł	PPTFPHSSYSVYIAENNPRGASI
		l l				FLVTAQDHDSEDNAQITYSLAE
		l				DTIOGAPVSSYVSINSDTGVLY
		1				ALQSFDYEQLRELQLRVTAHDS
		1		l	İ	GDPPLSSNMSLSLFVLDQNDNP
		1		1	İ	PEILYPALPTDGSTGMELAPRSA
		1				EPGYLVTKVVAVDKDSGQNA
	i	1				WLSYLLLKASEPGLFAVGLYTG
		1			1	EVRTARALLDRDALKQSLVVA
		1				VODHGOPPLSATVTLTVAVAD
1		l				SIPEVLADLGSLEPSDGPYNYDL
23812	54180	В	23950	165	335	SILE VENDESCRET SPOT TIVIDE
23812	54180	A	23950	217	859	GSAMAQQRALPQSKETLLQSY
23813	J#181	l^	23931	21/	0.55	NKRLKDDIKSIMDNFTEIIKTAK
		ŀ			İ	IEDETQVSRAT\QGEQDNYEMH
						VRAANIVRAGESLMKLVSDLK
						QFLIL\NDFPSVNEAIDQRNQQL
		1	1			RTLQEECDRKLITLRDEISIDLY
		l				ELEEEYYSSSPGPSLVSQQLSEQ
1	1	l	1			QLLSSAWTAWPKVTLSSLAPGP
1		ŀ				QYSVNGPAAIIAHPGTCASAGIS
		L_				SSLLPEPVLLVPVF
23814	54182	Α	23952	1	1263	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino aeid	*=Stop eodon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	İ			sequence		
23815	54183	ΙΛ	23953	70	2530	L PGVSRLVSKLRRARGMGSGAG
23813	34183	A	25955	l' ⁰	2550	
						ELGRAERLPELFLLTLFCPAL
						CEQIRYRIPEEMPKGSVVGNLA
	İ					TDLGFSVQELPTRKLRVSSEKP
	į.					YFTVSAESAELLVSSRLYREEIC
	İ					GKKPACALEFEAVAENPLNFY
	1		l			HVNVEIEDINDHTPKFTQNSFEL
						QISESAQPGTRFILGSAHDADIG
						SNTLQNYQLSPSDHFSLINKEKS
	1	l				DGSKYPEMVLKTPLDREKQKS
	ļ.					YHLTLTALDFGAPPLSSTAQIH
						VLVTDANDNAPVFSQDVYRVS
	l					LSENVYPGTTVLQVTATDQDE
	ì	l				GVNAEITFSFSEASQITQFDLNS
	į.					NTGEITVLNTLDFEEVKEYSIVL
						EARDGGGMIAQCTVEVEVIDE
	Į.	1				NDNAPEVIFQSLPNLIMEDAEL
		1				GTHIALLKVRDKDSRHNGEVT
		1				CKLEGDVPFKILTSSRNTYKLV
1		1			1	TDAVLDREONPEYNITYTATOR
		1				GKPPLSSSSSITLHIGDVNDNAP
						VFSQSSYIVHVAENNPPGASISQ
	1	ł				VRASDPDLGPNGQVSYCIMAS
	1	ł				DLEQRELSSYVSISAESGVVFA
	1	1				ORAFDHEOLRAFELTLOARDO
		1				GSPALSANVSLRVLVDDRNDN
						APRVLYPALGPDGSALFDMVP
	l .					HAAEPGYLVTKVVAVDADSGH
	l .	ŀ		1		NAWLSYHVLOASEPGLFSLGLR
					İ	TGEVRTARALGDRDAVRORLL
						VAVRDGGOPPLSATATLHLVF
		ł				ADSLOEVLPDITDRPDPSDLOA
23816	54184	С	23954	75	149	The state of the s
23817	54185	A	23955	442	618	CORLRSWVSOPFSFSFFPSFPTF*
23017	34103	1^	23733	1772	010	NILCLFVYLEVEFPSCCPGWSN
						NWAKSSGKALEAN
22010	64106	 -	22056		245	INWARSSURALEAN
23818	54186	A	23956	1	245	l

SEQ ID	SEQ ID NO:		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, ≔possible nucleotide insertion)
23819	54187	Α	23957	30	1079	APCASFLTPLAAASVGKVWLG
						LRGAAPMASVLSYESLVHAVA
						GAVGSVTAMT\VVFPLDTASIF
					į.	DFSVDF\KRKSKTTHMVVLEIIK
				1		EEGLLAPYRGWFPVISSLCCSNF
						VYFYTFNSLKALWVKGQHSTT
						GKDLVVGFVAGVVNVLLTTPL
						WVVNTRLKLQGAQFRNEDIVP
						TNYKGIIDAFHQIIRDEGISALW
						NGTFPSLLLVFNPAIQFMFYEGL
			l	1		KRQLLTFAFQLSSLDVFIIGAVA
			ŀ			KAIATTVTYPLQTVPLFPQFGR
	ľ		l			HRLNPENRTLGSLRNILYLLLQI
						FRTPSEREPFYTSVTLSVLSSVA
						YPMMPLSNITSAYRIGVLEKGG
			l	ĺ		GSGKSDNSDDDDDDDDDGGV
23820	54188	A	23958	1375	2451	VPRVPFAINVLTHSRQRYRRNL
						RFYADDPTIRV/GGPTYHWVRE\
						AEEERVVDNRMHDRFCELRTA
						AGHPVEGGRPLVIKGITHPAET
						ALQRFKFLMYQVVASDLDGTL
			l			LSPDHTLSPYAKETLKLLTARGI
						NFVFATGRHHVDVGQIRDNLEI
						KSYMITSNGARVHDLDGNLIFA
						HNLDRDIASDLFGVVNDNPDIIT
						NVYRDDEWFMNRHRPEEMRFF
						KEAVFQYALYEPGLLEPEGETK
						ALLTPGKLKNFLVEKMRTLGT
	1					AACPPYHIAFVIGGTSAETNLKT
	1					VKLASAHYYDELPTEGNEHGQ
		l				GSAMSSWNRNCWKRPRNSVLA
		l				RSLAVNTSRTTFALSVCHVTAH
		1				PARSAWASPVR
23821	54189	В	23959	1	1386	
23822	54190	Λ	23960	1375	1609	VPRVPFAINVLTHSRQRYRRNL
						RFYADDPTIRV/GGPTYHWVRE
				1		S\EEERVVDNRMHDRFCELRTA
						AGHPVEGGRPLVIK
23823	54191	Α	23961	433	741	
23824	54192	Α	23962	12	224	
23825	54193	Α	23963	3	2053	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /-possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1				sequence		
23826	54194	A	23964	1	2408	MGGADIAGPRREHNAKHPTVG
	1		1			LGSAAQTTCLPGPKQKGDPTRH
			i			QPVDKRNDVGPRQHFWCWRV
			l .			EIHKEKRMKTGPLNESELEWLD
						DILTKYNTDHAILDVAELDGLL
						TAVLSSPQEIEPEQWLVAVWG
1						GADYVPRWASEKEMTRFMNL
			l			AFOHMADTAERLNEFPEOFEPL
i						FGLREVDGSELTIVVPPGGRTA
	İ					TNALSCGSLFGEETSPRGPSADS
1						TSOSIGGGSPHDCAKGIALVAA
	İ					NGGDIRDYEGVDRSAKPOLPMI
			ļ			AINTTAGTASEMTRFCIITDEAR
						HIKMAIVDKHVTPLLSVNDSSL
1						
	ŀ					MIGMPKSLTAATGMDALTHAI
						EAYVSIAATPITDACALKAVTM
	1					IAGISTFFASSRMALMQASAPSA
						SFLPVTFTPIAAAQSRRRAAAT
			1			GYRDRWLCHGDVLLQICANTQ
						DTVIHALRDIIKHNSCPDDCSVQ
į.						HDAVIDDGCDFIDYLGGSETVV
1						ALNTPRADDYHPVMVLCAKTD
1						KAHRNVIHTSKLCFITLTPEISQ
						RSHGKSSKRVFSCFPVFLLLKH
						RRIVAQKYAELAYDLFHLGFD
						VLIIDHRGQGRSGRLLADPHLG
						HVNRFNDYVDDLAAFWOOEV
						QPGPWRKRYILAHSMGGAISTL
	i		1			FLORHPGVCD\AIALTAPMFGIV
						IRMPSFMAROILNWAEAHPRFR
1						DGYAIGTGRWRALPFAINVLTH
			1			SRORYRRNLRFYADDPTIRVGG
22027	64106	<u>_</u>	22077	,	22.62	SKOKTIGGER TADDITIKYGO
23827	54195	В	23965	14602	3357	CICDODDI BODI DA A TOTOLO
23828	54196	A	23966	14682	14931	EIGPGPRPLPSPLP*ATSTSVLAA
			1			SGRPERTR\HAGIKIVLEDIFTL
						WRQVETKVRAKIRKMKVTTKV
	l					NRHDKINGKRKTAKEQSPLLQE
1	l	1	1			SLFATGDVSHNLLRALDVGLLA
1		1	1			NLSALAELDISNNKISTLEEGIF
			1			ANLFNLSEINLSGNPFECDCGL
			l			AWLPRWAEEQQVRV/AAARGS
1	1	1	1			HVCWAWLPGWPASAWHPLAG
			1			QWLW*GVCRLPP*QQLRHRGS
1	l	1				SVLFSCPRRPA/CSQRPAAPSAS
						PPARASQPSPSRA
	L	L			L	TITIOTOGISTORY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
23829	54197	A	23967	1	714	MVKLSIVLTPRFLSHDOGOLTK
25025	15.1.57	ľ.	120,00	1		ELOOHVKSVTCPCEYLRKVINT
		1				LADHRHRGTDFGGSPWLLIITV
		1			l	FLRSYKFAISLCTSYLCVSFLKTI
		l				FPSONGHDGSTDVOORARRSN
		l			İ	RRRQEGIKIVLEDIFTLWRQVET
		l			ĺ	KVRAKICKMKVTTKVNRHDKI
		l				
		l			ł	NGKRKTAKEHLRKLSMKEREH
	1	1			i	GEKERQVSEAEENGKLDMKEI
	1	ı	l			HTY/MITPSARKPLCNRVRMAA
		_				AEHRHSSGLPYWPYLTAETL
23830	54198	Α	23968	111	216	ALLFPISTWSTPLGTQQAPQCV*
						PTPMPMRPVPS
2383I	54199	A	23969	I	4710	
23832	54200	A	23970	3544	3723	RSHRSPPRCPPSRAPAPRRSAAA
ŀ	i	l			İ	PPRR*PPRAPAAPRPGPP/CMAR
1	ł	ı				PPPARPTAGSRVPPL
23833	54201	A	23971	3	1119	EFGTSHRHRGTDFGGSPWLLIIT
	1	ı				VFLRSYKFAISLCTTYLWVINTL
		i			i	ADHRHRGTDFGGSPWLLIITVF
		ı		ŀ	1	LRSYKFAISLCTTYLC\VSFLKTI
	l					FPSQNGHDGSTDVQQRARRSN
		l				CRROEGIKIVL\EDIFTLWROVE
	1	l				TKVRAKIRKMKVTTK\ATRLTK
		l				IKERRKTAODHWRKLSMKERE
		l				HREEEROVSEAEENGKLDMKEI
		l				HTYMEMFORAQALR\RRAED\Y
	l	l				YRCKITPSARKPLCNRVRMAAA
	l	1				EHRHSSGLPYWPYLTAETLKNR
		1				MGHQPPPPTQQHSIIDNSLSLKT
		l			1	PPECLLTPLPPSALPSADDNLKT
		1				PAECLLYPLPPSADDNLKTPPEC
	l			1		LLTPLPPSAPPSADDNLKTPPKC
		1				VCSLPFHPORMIISRN
23834	54202	A	23972	1	1149	VCSLFFRIQKWIISKN
23034	34202	lv.	23912	l'	1149	

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
110.	sequence	nou.	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		l		sequence		ĺ.
		<u> </u>				
23835	54203	A	23973	93	1165	HRRSAAEAPAAAVMPFPFGKS
						HKSPADIVKNLKESMAVLEKQ
1			l			DISDKKAEKATEEVSKNLVAM
l						KEILYGTNEKDPQTEAGAQLAQ
1	1	l				ELYNSGLLITLVADLQLIDFEGK
	1					KDVAQIFNNILRRQIGT\RTPTV
l	1					EYICTQ\QNI\LFMLLKGYESP\EI
ĺ	1					ALN\CGIMLRECIRHEPLGKIIL
i	i					WSEQFYDFFRYVEMSTFDIASD
						A\FATFKGLLTRHKLLSAEFLEQ
	1					HYDRFFSEYEKLLHSENYVTKR
			l	1		QSLKLLGELLLDRHNFTIMTKY
			l			ISKPV\NLK\LMMNLLRDK\$RNI
			1			QFEAFHVFKA\FVANPNKTQPIL
	1					DILLKNQAKLIEFLSKFQNDRTE
						DEQFNDEKTYLVKQIRDLKRPA
						QQEA
23836	54204	Α	23974	3	158	
23837	54205	A	23975	3	1071	GVRASGRAVPRAVFAGMKRPC
						EETTSESDMDETIDVGSENNYS
ŀ						GQSTSS\VIRLNSPTTTSQIMAR
						KKRRGIIEKR\RRDRINNSLSELR
l			İ			RLVPTAFEKQGSAKLEKAEILQ
l	1				l	MTVDHLKMLQATGGKGYFD\A
l	1					HALAMDFMSIGF\RECLTEAAR
						YLSSVEGLDSSDPLRVRLVSHL
					ľ	STCATQREAAAMTSSMAHHHH
	1					PLHPHHWAAAFHHLPAALLQP
						NGLHASESTPCRLSTTSEVPPAH
						GSALLTATFAHADSALRMP\ST
l	1	l	1	1		GSVAPCVPLLSTSLLSLSATV\H
l	1	l	1			AAAATATAAAHSFPLSFAGAFP
l	1	1	1			MLPPNAAAAVAAATAISPPLSV
	1					SATSSPQQTSSGTNNKPYRPWG
l	1		l	1		TEVGAF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23838	54206	A	23976	1	1416	MPGNLTRKESSSPLTDRHCHTS
						SQFSHRFNAISPFPPAELGCDHV
		l		l		ASAAADPLGVPGVATQWAGGL
		l		l		SAALAAPRRPHSCRRLOVLGAL
		l				GILKSAGRVNFRCFGPLSIRGKP
		l		ł		KTPLWEQEKPKLQTGSAGSQQ
		l				KRVMLKPSRGADRVLGRGSQR
						PAETDRETTDLRQRKRMKEGK
						ERTLRAARGETRRSNGNLARO
			1			TVPEGCRSQTQALQVVRNART
		1				PPCIPNLDRPPEACSSGSTGGM
				i		MPLRVAPSPVASGESSHLSSGR
		l		l		GDRGLVRCKVEELRFQPVAFPE
				1		FIPPPTSPRADSESGDWCCTVLA
						NPSKTGSGGWEQRKTEC\PQGR
		l				TONPOORGS/RNLVLLTAGOES
						TPRRSQLTRSPAAERRTEEQGE
		1				DRKNRALGCNLFPRVPRHARD
		1				SORALRASTOAPKKSTFAFPGI
						HSGAVCSKRRICGVQGEPFPLPF
						ASQCSSCLKILPSPGPTKELTSY
						LLHSGCGFLDKLRV
23839	54207	Α	23977	3	139	
23840	54208	A	23978	141	734	GSQLRRPAFGVRVALEESKGLI
						RKDSCRGRACAGANACARRRR
l		1				PPTMSGAAWRGTPLALVRSSRT
t		1				RGVWAWQWSATRSPSDSCFPG
		1				VARDKEDGEPKTKHLREEEEE
		1				GEKHRELRLRNYVPEDEDLKK
						RRVPQAKPVAVEEKVKEQLEA
			1			A\SPEPVIEEVDLANLAPRKPDW
l				1		DLKRDVAKKLEKLKKRTQRAI
						GRLDP
23841	54209	Α	23979	3	585	GTVGGVCGPRRSSSASPQEQDQ
					1	DRRKDWGHVELLEVLQARALF
		1				TAVPLSEEDRATVLKAVHKVK
					1	PTFSLPLQLPPPVNTSKLLRDVY
		l				AKDGRVSYPKLHLPLKTLQCLF
		l				EKQLHVELASRMCVVSVEKPT
		1				LPSKEVTHARKT\LKTLRDQWE
1						KALCRELRETKNRLERQVYEG
1	I	1			I	RFSLYPFLCLLDECEVVRMLLQ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	eodon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23842	54210	Α	23980	1	3778	MRRRGPVGWPGAACVMSALC
		1				WGRGAAGLKRALRPCGRPGLP
		1				GKEGTAGGVCGPRRSSSASPQE
		l				QDQDRRKDWGHVELLEVLQA
		l				RVRQLQAESVSEVVVNRVDVA
		l				RLPECGSGDGSLQPPRKVQMG
		1		1		AKDATPVPCGRWAKILEKDKR
		1				TQQMRMQRLKAKLQMPFQSG
		ł	l			EFKALTRRLQVEPRLLSKQ\MG
		l				GCLEDCTRQAPESPWEEQLARL
		1				LQEAPGKLSLDVEQAPSGQHSQ
						AQLSGQQQRLLAFFKCCLLTDQ
23843	54211	Α	23981	101	805	HMYTSMKPSSQPRSWISTPHPK
						ACSSLSFPLSLDNWMNINLREVI
						CTSRDGDKFWRMPECYIRGSTI
			1			KYLRIPDEIIDMVKEEVVAKPR
		1				GRGGLQQQKQQKGRGMGGAG
		1				RVGSPLSSG\VPGAGRQPHVAS
		l				LLLGVALAALPRRLPNLPGPPS
		1				CPLAMNRGPGRGAPRHGTQPSS
		1		1	l	QCASWRSSGHIKSPPQAPVSPM
	1	l				RGNGGAHLSRPLATTLGTLRGS
				ļ. <u>.</u> .		IQSQGRPPPPEMMNWH
23844	54212	A	23982	133	1069	RIAWGQDRSCLQGGPFATYVP
						GLILCSLQGWYGVQAGQDLQG
	ì			1	İ	LPRSQGWARRPGGKLITCSPSP
					1	QDGDKFWRMPECYIRGSTIKYL
		i				RIPDEIIDMVKEE\VVAKGR\GR
		1				EGPCQQAESQQK/GVRGMGGA\
		1		1		GRCVFGGRGRGMGSPITG\RCH
		ł		1		PDKKPGRQAGQTVSAHPDRLL RPLLARVGDFRSTVSERNLLOK
		l		1		GRGRPPRGEPPFRVLVTPSLFKV
	1	1		1		
	İ	1		1		KAPFFLPKNRQILRVRKCLLCCF FGFCETGPSOKLGWNPLERVLC
		1			1	LKAIFSSPFASGFSRASHFPRRG
	i	1			l	LFLHFLGSLSEASNOTGRFLILK
23845	54213	A	23983	1	326	PVSLECYTVPPEDNLALLQLYF
23043	54213	۱^	23703	1.	1320	RTLVTGA/LRPRWCPVLYAVTL
	l	1		1	1	WPHVNSFIFSQRPRIVDIDGKM
		1				GTLSCIOPFNRCLARRRTLCPNG
	1	1		1		ROETKEOKACLPGSRLFELF
23846	54214	A	23984	2	341	NYL TREORACLE OSKLIELE
23840	134214	I ^A	23904	14	Dat	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
		!		 		
23847	54215	A	23985	114	712	DLISALMIWFHVIVIHTPYVCIYI
		l				PTLPLSTSFPLSQLPVSLECYTV
						PPEDNLALLQLYFRTLVTGALR
		1				PRWCPVLYAVAVAHVNSFIFSQ
						DPQSSVSYISIFRGRKGASN*AA
		ļ.				SGPAGEEVSVKQGQWWVPAL
		l				GSVPPLPVLLL\DEVKAARRSM
	İ	ŀ				LQKTWLLADEGLRQHLLHYKL
						PNSTLPEGFELYSQMPPLRQHY
23848	54216	Α	23986	357	2771	FESSCPPVGSPVGAGPTPSLYKV
						PCATAMKLLRVLASAGRNIAA
		1				RLLSSFDLRSRLCRIIAEAPQEL
						ALPPEEAEMLSTEALRLWAVA
						ASYGQGGYLYRELYPVLMRAL
						OVVPRELSTHPPOPLSMORIASL
		l				LTLLTQLTLAAGSTPAETISDSA
	l	l				EASLSATPSLVTWTQVSGLQPL
		l				VEPCLRQTLKLLSRPEMWRAV
		l				GPVPAACLLFLGAYYQAWSQQ
		ı				PSSCPEDWLQDMERLSEELLLP
		l				LLSQPTLGSLWDSLRHCSLLCN
		l				PLSCVPALEAPPSLVSLGCSGGC
İ		l				PRLSLAGSASPFPFLTALLSLLN
1		1				TLAQIHKGLCGQLAAILAAPGL
		1				QNYFLQCVAPGAAPHLTPFSA
		ļ.			1	WALRHEYHLOYLALALAOKA
						AALOPLPATHAALYHGMALAL
	l					LSRLLPGSEYLTHELLLSCVFRL
		1				EFLPERTSGGPEAADFSDQLSL
		1		İ		GSSRVPRCGQGT\LLAQACQDL
ŀ			į.		1	PSIRNCYLTHCSPARASLLASQA
					l	LHRGELQRVPTLLLPMPTEPLL
		l	į .			PTDWPFLPLIRLYHRASNTPSG\
		l				LSPTDTMG\TAM\RVLQWVLVL
			i			ESWRPOALWAVPPAARLARLM
1		l			ŀ	,
1		1				CVFLVDSELFRESPVQHLV\AAL
1	l	1	ł			LAQLCQPQVLPNLNLDCGLRG
1					1	LTSFPDLYANFLDHYEAVSFGD
					1	HLFGAMVLLPLQRRFSVTLRLA
		l			1	LFGEHVGALRALSLPLTQLPVS
						LECYTVPPEDNLALLQLYFRTL
23849	54217	Α	23987	291	454	KAETGSLPYTLCKNQFKMD*RL
			1			TC*T*NHKNPRRKPR*YHSGHR
		_				HGQRLHDENT
23850	54218	Α	23988	1	613	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /-possible nuclcotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23851	54219	A	23989	807	1345	RCEERLLPLNLAVIALPSGLER
		1				WSFFAVYNGHPGSQVAKYCCE
l		l				HLLDHIADNQ/DFTGSAGAPSLE
		l				NVKNGIRTG/CPEPEVHDTERSE
		l				EDDHFIILACDSIWDVMGNEEL
		l				CDFVRSRLEVTDDPEKVYNEIV
		l				DTCLHKGSQDNMCVIF\LWFPN
						APKVSPEAAKKEAELGKYRLN
		l				SILSDQ
23852	54220	A	23990	344	636	SSLSLLDIWVGSKSLLL*IVPQ*T
		l				YVCMCLYSSMIYSPLGIYPVMG
	İ			i		WLGQMVFLVLDPSGIATLTST
	i	l				MVELVYSPTNSVKVFLFLRILSS
		l				TCCFLTF
23853	54221	Α	23991	l	2793	
23854	54222	A	23992	18	409	SSVVEFVDPRVAPALLPQLPLES
						SSHGSVLPTMAVCLDKGHKMT
		l				KNMSKPRHSCRRGRLTKH\KFM
		l				WDMIREVCGFAPYERHAMVLL
		l		İ		KVSKDKRALKFIKKRVGTHIRT
	1	1				KRRQEELSNVPAIMRKADAKK
						D
23855	54223	A	23993	1	609	MHPRGPPPPAQEESHTDESPGID
l		1				HQSLARRAGRVGRTGGFSQLL
		1				AQTNTSMLSSSPASCTSPSPDGE
		l				NPCKKVHWASGRRRTSSTDSES
l						KSHPDSSKIPRSRRPSRLTVKYD
		1				RGQLQRWLEMEQWVDAQVQE
	1	l				LFQDQATPSEPEIDLEALMDLST
						EEQKTQLEAILGNCP\RPTEAFIS
		I				ELLSQLKETPETQPASEISLRDH
23856	54224	Α	23994	I	204	
23857	54225	A	23995	1	418	MWNRLCNWVTSIGWKSLEGSE
l						EDRKSSQWGTLPSGAVRRGPPS
	1					SRSKNGRSNSSLHPVPGKATGT
	1			1		QLEPMRAVMGDESCKATGTEL
	1			1		LHALGAHPLHRLLKQHSHMMF
	1	1		1		QKVQVEPLRK\EPHHWQRL*PF
	1					SSE*NHGPSR
23858	54226	A_	23996	104	253	

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		·
23859	54227	Α	23997	666	1436	WRPCFWLQAKKEAPAP\PKAK
23003						AKVKAL\KAKKAVLKDVHSHK
		1				KKKD\RKSPTFRG\PKTLCTR\KP
		l				PKCLQKTAPRKNKLDHCAITKF
		l				PLTTKSTTKKIEDNNTLVFIVDV
		1				KANKHOFKOAVKKLYDTDVA
		l				KNITKRWDVTSIITSCYIHLRLA
		l				SRLALEDLLAGLMNCLKGTFW
		1				NEESGERVETRPESRFKQQSKLI
		1				YTIKNPGRKKYGAEEKRAFWC
		1				HTDLLLHDHDEQVNHFSSPSST
		l				FTKCKRKKPFLLREFVWIK
23860	54228	A	23998	2	301	FIRCERORFFEEREFVWIR
23861	54229	A	23999	1	242	MGMVRKQALERTEWALVCRQ
23001	J 122)	ľ`	23,777	ľ	1	AWSSRMGYSTTVSPTOCOYTC
						SRGLLDSOLOLYKCTOALPEIH
		l				SLGHVCPSLITVKYSHYVLAAA
				ł		TLFIWDNNESLRTGLLSVSTCK
						SGTRDFMRISGMVSLTSEVDNV
		l				EVFHYFPLHIQGKKFWAVGAI
	1	1				MTIFVISIYYEPGAHWHHMWK
						LLKVYSLESQKREEDISKRLPNF
		1				LKSSRRPWEDLILMLRKSPEMS
		1				KEYRASPLSLSAALRSKDRQSR
		1				HRCVTSTFHIREVRCCKSQGLP
		1				LCMHETAAHICP*TEGOOKGSG
		1				GLLYIP*TSPEIFSASESDPPMAC
23862	54230	Α	24000	2	948	CGTPTVVPGTGTSRRVRTATVE
				1		PTWKCRAAQQPPPPP/PPLWVK
			ŀ		ł	LPPPDTSTKPELQLEELDADSS
	i			l		WMPLHGSCPTGKLAVDARSSL
	•	l		l		FTEPAFPYWKKMKHTFQNILS
	ŀ	l				WSKFTNLTYSVTQLWLLVVMA
	İ	1				QTSYPHSLVGRKQVILHSSCLY
		l				SLDNKTSLAGDDHDDDDHDDN
		l				DSSPAEKGPPPITKSLDKADEQS
						HLRLAVLELGFYKPVNLLLVA
		1				MTGCLLQLMMKKEQCGTHGG
		l	i			EGSTLEGSRPDVPQDLFSPTLGF
		1				YGSTSLSADLGENVILKLYLPSF
						PLQESEPSAEHGENLWALYVGF
						EERRSEASLAKAP
23863	54231	Α	24001	265	513	
23864	54232	Α	24002	513	796	TGTAWTPPPPPLTTGAPCTPPPR
	1	1		I	1	CTARGRT/PGDSHLGGGPAATA
	1	l	1	1	1	GGPRTSPMSSGGPSAPGMRPPA
				1		SSPKRNTTSLLNSGLEPTFSFRIT
			L			FGFM

SEQ ID NO:	SEQ ID NO: of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *-Stop coden, /-possible nucleotide deletion, \-possible nucleotide insertion)
23865	54233	A	24003	1268	2097	TTELTGAISLSTLGDPSPFRQPS AQMGHLLSTSLSTPPTISSSYLPT QPAFATVTPSSSSVLPGLPQTSF SGMGPSAELMASTISPKQQLPQF SAAFGHQLSSHSGIPKDLQPSHS SIAPPTGFTVTGVKLQTFAVSVT VLKATCLELFVPPGGFVVSLAS GVKLQTVTPLHFSGNHPSAFYY YRLSFEMDKGKKRQFGVEKEK CGSLGGFVMLGNEBMIRKAV LLAWKLIPFLGIREAKNPGSENI RLATILEAPPPSWELWEQGPPG APYYQP
23866	54234	A	24004	3	1163	GEVTSWNRGGGLRIWRYFLEGD GVOKTMELLISRLTYCWLEGDG RVSWRLGLEIGSLPTPVPVRLCP ANKSPPMVCGTTGRSSNSSTST KRMCTCSSRAALIFILKGPEGG TLMEEIRALIKEPPESSLTPSTRG GHSEKAPSKNHQAATVNTTLIS RISHISHTHLLIHCTGKEQHEPY EDAATRSRPEADLSRHERLTCR HLDLGLPDMONRQCVGSGGL RPVGGLEGPRHWCLVRSSRLG VLPRRPQGCEDDAGGLQRQDC GQWRELVQLPTPPVCPGGASP HPKPLPDLPDPPDSSRCPCLLPG TCPAAPPAGSPSPSCSSPCCLLPG VPTPSVLECCTQVLLEDSKPQS HRNSAPHLQPGIWSHIPGSHPW CLVLAAEGSTSSV

SEQ ID NO:	of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23867	54235	A	24005		2502	MEARVERAYOKROVLFLCVFL GMSWAGAEPLRYFVAEETERG TFLTNLAKDLGLGVGELRARG TRINSDONMOILLISSLTGDLLL KEKLDREELGGPREPCVLPFQL LLEKPFQIFRAELWYRDINDHA SYFLDREISLKLIESTTGAAFLL ESAQDSDVGTNSLSNYTISPNA YPHINVHDSGGGINYPELVLNQ VLDREEIPEFSLTLTALDGGSPP RSGTALVRILVLDVNDNAPDFV RSLYKVQVPENSPVGSMVVSV SARDLDTGSNGEIAYAFSYATE RILKTFQNPTYGSLHKAQLDY EAIOTYTLTQAKDGGGI,SGKC TVVVDVTDINDNRPELLLSSLT FYTLVTIGAKDGGGI,SGKC TVVVDVTDINDNRPELLLSSLT FYTLVTEKPLDKERNTEVNITIT VTDLGTPRLKTEHNITVLVSDV NDNAPAFTQTSYTLEVRENNSP ALPIGSVSATRDBGSTAGNSP ALPIGSVSATRDBGSTAGNSP ALPIGSVSATREFRNNSP ALPIGSVSATREFRNNSP ALPIGSVSATREFSRNSP ALPIGSVSATREFFRNSP FALHIGSVSTAETNIQVTTSLLP PRNPHLPLASLVSINTDNGHLF RSLDYFMFHLPLASLVSINTDNGHLF RSLDYSSTAETNIQVTTSLLP PROPHLPLASLVSINTDNGHLF RSLDYSTAETNIQVTTSLLP PROPHLPLASLVSINTDNGHLF RARSLDYSALGFEFFRVGASDR GSTALSSEALVRVLVLDANDSS LFVLFPLQNGSAFCHEMPRAVA AEPGYLVTKNAWLS*QLLKAT EPGLEFGWWAHNGTDRLLSERD

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /-possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23868	54236	Α	24006	1	1960	MEARVERAVQKRQVLFLCVFL
		ı				GMSWAGAEPLRYFVAEETERG
						TFLTNLAKDLGLGVGELRARG
		l				TRIVSDQNMQILLLSSLTGDLLL
						NEKLDREELCGPREPCVLPFQL
			ł			LLEKPFQIFRAELWVRDINDHA
						SVFLDREISLKILESTTPGAAFLL
						ESAQDSDVGTNSLSNYTISPNA
						YFHINVHDSGEGNIYPELVLNQ
						VLDREEIPEFSLTLTALDGGSPP
						RSGTALVRILVLDVNDNAPDFV
			1			RSLYKVQVPENSPVGSMVVSV
						SARDLDTGSNGEIAYAFSYATE
						RILKTFQINPTSGSLHLKAQLDY
						EAIQTYTLTIQAKDGGGLSGKC
						TVVVDVTDINDNRPELLLSSLT
						SPIAENSPETVVAVFRIRDRDSG
	1		l			NNGKTVCSIQDDLPFILKPSVEN
						FYTLVTEKPLDRERNTEYNITIT
			1	ľ		VTDLGTPRLKTEHNITVLVSDV
			1			NDNAPAFTQTSYTLFVRENNSP
			1			ALPIGSVSATDRDSGTNAQVIY
						SLLPSQDPHLPLASLVSINADNG
			İ			HLFALSPWTTMPLRAFEFRNNS
			ł			PALHIGSVSTAETNIQVTYSLLP
			Į.			PRNPHLPLASLVSINTDNGHLF
			l			ALRSLDYEALQEFEFRVGASDR
		l				GSTALSSEALVRVLVLDANDSS
						LFVLFPLQNGSAPCTELMPRA\A
23869	54237	Α	24007	1	966	MGHHLMIDKLVALGGLYYAIQ
					ľ	RHYATKCSVLKNDQILVIGLFM
			ŀ			IQNVIYRKHFANPLSALFLQQGI
						ELFAAIAEIHITTVAERNHAITQI
				1		RLEAQTIFDSLKEWHNAIRKSL
						DYEALQAFEFRVGATDRGFPAL
				ĺ		SSEALVRVLVLDANDNSPFVLY
	i					PLQNGSAPCTELVPRAAEPGYL
	l					VTKVVAVDGDSGQNAWLSYQ
						LLKATEPGLFGVWAHNGEVRT
						ARLLSERDVAKHRLVVLVKDN
	1					GEPPRSATATLQVLLVDGFSQP
	l					YLPLPRAAPAQAQADSLTVYL
						VVALASVSSLFLFSVLLFVAVR\
			L			LCRRSRAAPVGRCSVL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nuclcotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			i	sequence		
		<u> </u>		ļ		
23870	54238	A	24008	[1	2673	MEVMGNMGSKILITGITSTKRK
		1				VALLAKKRERKIISQRKDVTDS
1						RAKRQSEEKSIGKETVVIGIGLI
1					i	SKDSAAAICESRWRLSESHFER
						LIQRRMEARVERAVQKRQVLF
	1					LCVFLGMSWAGAEPLRYFVAE
1		l				ETERGTFLTNLAKDLGLGVGEL
						RARGTRIVSDQNMQILLLSSLT
1						GDLLLNEKLDREELCGPREPCV
						LPFQLLLEKPFQIFRAELWVRDI
	1					NDHAPVFLDREISLKILESTTPG
	1		i			AAFLLESAODSDVGTNSLSNYT
						ISPNAYFHINVHDSGEGNIYPEL
1						VLNOVLDREEIPEFSLTLTALDG
l	1					GSPPRSGTALVRILVLDVNDNA
	1	i			j	PDFVRSLYKVQVPENSPVGSM
	l					
1						VVSVSARDLDTGSNGEIAYAFS
		1				YATERILKTFQINPTSGSLHLKA
						QLDYEAIQTYTLTIQAKDGGGL
	İ	1				SGKCTVVVDVTDINDNRPELLL
	i	1				SSLTSPIAENSPETVVAVFRIRD
						RDSGNNGKTVCSIQDDVPFILK
	1					PSVENFYTLVTEKPLDRERNTE
	1					YNITITVTDLGTPRLKTEHNITV
					1	LVSDVNDNAPAFTQTSYTLFVR
						ENNSPALPIGSVSATDRDSGTN
						AQVIYSLLPSQDPHLPLASLVSI
ŀ				1	1	NADNGHLFALRSLDYEALQAF
1		ļ.		1		EFRVGATDRGSPALSSEALVRV
1	Į.					LVLDANDNL\PFVLYPLQNGSA
}						PCTELVPRAAEPGYLVTKVVA
Í		ľ	ŀ			VDGDSGQNAWLSYQLLKATEP
23871	54239	В	24009	1	993	T D G D G Q T T T T T T T T T T T T T T T T T T
23872	54240	В	24010	i	4056	
23873	54241	В	24011	i	906	
23874	54242	В	24012	li .	4318	
23875	54243	В	24013	41	1935	
23876	54244	Ā	24014	3	328	GWOSGPVLGGLCGTAGOSCGD
200.0		'		ľ		GDGIWWAQAAQAGDPAHLRL
1	1		Į.			LTHGQPSSVPEAPVRAHQPALP
						QPRSAAARECGR/AGRSTTNSL
	i		l			RKRVASESYMIVAPPMPSSWSS
	l					
22275	-1015	l	24015	114		SG
23877	54245	A	24015	114	567	VKKPHSAEWVPLWGLQPGWR
		1	ĺ	ł		QPFRCNSGSEKELRRPVSNSQEI
				1		LSPVTMRKPAVSVCCDLTLREA
		1	l	1		APPFQRHERTYTGEKPYKCKQC
1		1		1		GKAFSGFCYTQILEKAQTGKQP
		1	l	I		YE/AHERTHWKQNHDCTQFGK
1		1	l	1		AFSFSSSLCIHEKIHTGEKLYE
	L					

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nueleotide	Nucleotide location of last	Amino aeid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequen c e	deletion, \=possible nucleotide insertion)
23878	54246	Α	24016	170	491	PAQHQRGLWQSPTHQALTLML
						SVHREDWKPVLTINSIIYGLOYL
						FLESKNRWLGLERLGFCPSGLIT
		l				PPTCTGHRSP/NPEDPLNKEAAE
		1				VLQNNRRLFEQNVQRSMR
23879	54247	A	24017	1	1062	
23880	54248	Α	24018	2	700	RRPTRPGGGSGGGGRMIKLFS
						LKQQKKEEESAGGTKGSSKKA
						SAAQLR\IQKDINELNLPKTCDIS
ŀ						FSRSQTTLLNFK\LVICPDEGFY
i		1	İ			KSGKFVFSFK\VGQGYPHDPPK
	l	1	i			VKCETMVYHPNIDLEGNV\CLN
		l				ILRGDWEASPLRLNSIIYGLQY\
1			ľ			LFL\EPNPEDP\LNKEAAEV\LQN
						NRR/VCFEQNVQRSMRGGLHPA
						PPTFERCLKIGLAHTHPPPRPQA
				1		LASPAKYLLGAMG
23881	54249	Α	24019	302	884	PMAPLSSQQPTLFPNTLTQAQQ
						VQLTKGLRKWMVQHGGDGW
						VVEENRTTVPGAPSQTCFVTSF/
						EVNAPAIQSWCCKKQVLDLEE
						EGLWPELLDSGRIEICVSDWWG
		1				ARHDSGCMYRLLVQLLDANQT
ŀ		1		1		VLDKFSAVPDPIPQWEQQCLPF
						TFTHVFFQHSRWGVRFCVFSEH
						RGPGTHSSGAGPLLSPCEQLQC
23882	54250	A	24020	274	866	SSLLNLPSSWDYGWVSYEQAN
	1					CKGEQFVFEKGEYAHWDWWT
						SSQRTDFSSLR/RPSKVDSQEPK
l						V\YLYENPNFTGKKMEIIDDD\V
İ						PSFHA\HGYQEKVSSVWVRSG
		l				MWVG\YKYPGYRGLQYLLKKG
		1				DYKDSSNF\GARHPQV\QSVRCI
		1				HDMQGHQCGAFHPLGTQEEEE
l		1				AGQKEQMEQVPFWLARGDICE
		L				LSGKKSVI
23883	54251	A	24021	179	374	AASIGRSMTEKAHAGPRPGRN*
1		1			1	ADAPSSSDTLFPPDALLSNQAG
						RLACWLFSYGPQTKNDFHIFK
23884	54252	Α	24022	1	141	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /-possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
23885	54253	Α	24023	Īī	506	PHSPLWKTLCSPSSGVCCGFVT
						LFCNDRFACLFQLTPPPSPAWM
		1	ŀ			TAAASLLASLLPALPPLNLQDA
						KFVEERRKQLQNYLRTVMNKV
		l				IQMVPEFAASPKKETLIQLMPFF
						VMGLQVGMYGLFHFGVMMDQ
						PQPSLTRADESPCDRVLSPCAA
			ŀ			HRKQRDCCLSWKAEPE*FLVK*
						KTLCSPSSGVCCGFVTLFCNDR
					<u> </u>	FACLFQLTPPPSPAWMTAAASL
						LASLLPALPPLNLQDAKFVEER
		ĺ				RKQLQNYLRTVMNKVIQMVPE
						FAASPKKETLIQLMPFFVMGLQ
				1		VGMYGLFHFGVMMDOPOPSLT
				l		RADESPCDRVLSPCAAHRKOR
						DCCLSWKAEPE
23886	54254	Α	24024	1	396	
23887	54255	Α	24025	1	1176	MVPGDLSQTSEDQSLSDFEISN
						RALINVWIPSVFLRGKAANAFH
						VYQQSIAGVSGIARKRALIPFGK
						STFSYVCITNIKLTDALPSWDYT
	ĺ			1		CHFNLPKAQKPGEPTKCILTME
			ł	1		WLALLQGDPGSWIQKEFIVGKR
			ŀ			GGKMDNILWDQRDVQPPGLRP
	l			1		HQLLHSGGGPLISFRNGGPILV
	ĺ		l			WELRRKGASGKEIVGSRYFRES
					1	DEVLVWKNPKTLQKEDEEVNS
		l				THPMHTYPCTHCQVFCVTAEII
		1				NIYLQACIVLEMSMLPPGERPLF
						LLRLTTVLEVTVFDSQWSRPYH
		ĺ				VSADPSAWKHHHHYHHHHHY
		l				QLSSSSPTITIINHPSPS/LSHASPS
	İ		l			P/CHPSPSPAVHHHHQPSITITSR
	1	1		l		PSPSPAVHHHHQPSITIIIRPSSITI
						INHPSQASSIHHHHQPSIPFIVSH
23888	54256	Α	24026	1349	2125	
23889	54257	Α	24027	670	972	KTDEWNIYRRYTEFRSLHHKLQ
		1				NKYPQVRAYNFPPKKAIGNKD
1		1		l		AKFVEERRKQLQNYLRSVMNK
		l				VIQMVPPSSLPAPKKETLIQ\LM
1		l	l	I		PLLRSTSTPPRRAL
23890	54258	A	24028	1	699	

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			l	sequence		
23891	54259	Α	24029	221	864	VDCPRSFSAVVVWQGRFWLLS
İ						FIPGGFLGMTLFFASGMNWTVT
			l			LLGFLAGNLEVEPSDTIENVKA
			l			KIQDKEG/IPPDQQRLIFAGKQL
				ŀ		EDGGTLSDYNIQKESTLHLVLR
						LRGGMKIFVKTLTGKTITLEVE
	ŀ	1	ł	i		PSDTIENVKAKIQDEEGIPPDQQ
ł		ĺ	ł			RLIFAGKKLEDGRTLSEAIFLDV
						GLLLSCRTVSSCGAFILGWSVVI
00000	54050	<u> </u>	0.4020		064	LIPVISTRDDTSTTK
23892	54260	Α	24030	2	864	PSFRLAWPEGRLALLGELVCVP VVGRGMENGRILVSADRSGGHI
						CLKTLT\GKTIT\LEVEPSDTIEN
						VKAKIODKEG\IPPDOORLIFAG
						KQLEDG\RTLSDYNIQKEST\LH
			ŀ			LVL\RLRG\GMQIFP*RP*PAKTI
						TLEV\EPSSHHRKF*RARFOGKE
			İ			GIP\PDQQRLIF\AGKQL\EDGRT
1						LSDYNIOKESTLHLVLRLRGGM
l						OIFVKTLTGKTITLEVEPSDTIEN
1						VKAKIODKEGIPPDOORLIFAG
						KQLEDGRTLSDYNIQKE\STLHL
i						VLRLRGGC*FFSHGIRSAQ
23893	54261	Α	24031	1	124	
23894	54262	Α	24032	97	296	
23895	54263	A	24033	71	269	AAGYRPCPRHPGARRTAATRR
						CHPSPAVARPAGCSGPP*CYAL
1						SPDPARTRRGDPQTPPCRLPRA
22005		<u>_</u>	21021	201		G STRLGLPKCWDYRREPPRPAKS
23896	54264	Α	24034	206	667	SFLTAQACEAQKTGSATRRKD
						CFLOPTFWLLPFLVLEKGAPSA
l .						CRGKRYVRAVWLSLSTGAILOP
						YPKLOVSNWL*CARDOEMPEG
1						CL*OSCOTLSLLVLPPKEKKDV
1						*EISPWTPAPPEHQYTVRTFQSP
23897	54265	A	24035	3	190	GTFKNDLILLISKSYY*APLPSD
				Ī		FSHSDTSS*KPSKSLKTYPSPPFF
			l			SAKTSLLQCPRNILY
23898	54266	Α	24036	197	474	SPLSHRGASELLSSLHLRSSCSS
			l			PKMAETPEAGFQIVHSASPQPG
	İ		l			TAPGNPWAAGAGG*HCEPGQL
1						AGPYRAASFFGQAWPPHPRAK
						LGVSR
23899	54267	Α	24037	164	323	GQEIKTILANMVIPRLY*KYKKI
			l			SWAWWHAPVVPATWEAEAGE
L			L			LLEPGRQRLQ

SEQ ID	SEQ ID NO:					Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540.217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		03/340,217	sequence	or pepulae sequence	decision, possible nacional insertion,
		<u> </u>				
23900	54268	Α	24038	582	1166	HCEPGQLAGHLGQPLSSGRLGF
				1		LMASHKLSNNYYRTRDGRRES
1		l			1	VPPSIIMWSQKALVSG\KPAERV
						YGDANHTVGNMCASRLGGYL
ł		1				GFSIRIVVFIWYLGLSVVVNML
ŀ						VGRQCDWQGLVYGQPELLVFS
						GNMPGEVARGKHRNEKAPLSM
		1			!	RPASQCRVFCVLKGCSERSRHK
		l				EQTPDTTIFKNCNTHCEGPWLH
23901	54269	Α	24039	1214	3878	
23902	54270	В	24040	62	1242	
23903	54271	Λ	24041	74	408	
23904	54272	A	24042	790	1877	DEIVDPGSMKWSYKSQLNYKT
l						KRPSVFVTALTHRPDILILGTLE
						SGHSRNLTCSVPWACKQGTPP
						MISWIGASVSSPGPTTARSSVLT
		l				\LLPQSPQDHGTS\LTCQCT\LPG
1		1				TGVTTTSTVRLDEAYPA\WNLT
		1				MTVFQGEGTASTTLRNGSALSV
l		1				LEGQSLHLVCAVDSNPPARLS
ł		1				WTWGSLTLSPSQSSNLGVLELP
						RVHVKDEGEFTCRAQNPLGSQ
		l				HISLSLSLQNEYTGKMRPISGV
						MLGAFGGAGATALGFLSFCIIF
		l				VGVRSCRKKTARPTMG\VGDT
						GMEDT\NAVRGSA\SQGTPELN
1						PRQKEQAPQNHGSANALANPP
į.	Ì					RPEEGEIQYASLSFHKARPQYP
1						QEQEAIGYEYSEINIPK
23905	54273	Α	24043	183	1176	VHQLVFPRQHQRHSVFPKANSS
						WIHQLFPQ\QHSYGPSAVSAGQ
						HQLMVHSLFPQATPAHASISLV
						FRQGTHQLHRVHQRRIPSAQHQ
	1	l				RHRLHQRRLPSAQHQRHRDSIS
	1					VCFPFRAHTRPHCDSIQRQHQL
						MRSIGLFSSGKTDPTVTLLKEA
	1					AAEHEICLCSRHPVNLRETKAH
		l				LGNAAHA VFPRGNQKKPEMCP
	ł					YLPTHKLVVPTPPLIPLEAALRN
						IAHSRSIPPPKKFSPTPDTSDDYL
						LVFNSFPYSKLTTPKTGQGENL
1	1	1			1	GSTRPIPCNGPPTRGPHNGAKY
		l				PGDVSHPVGVGTLWANRLLTG
		L	L			QYEENGKSFFLLICTIATFQVNA
23906	54274	С	24044	178	342	
23907	54275	С	24045	86	283	
23908	54276	С	24046	91	177	
23909	54277	С	24047	H	169	
23910	54278	С	24048	209	673	
23911	54279	С	24049	164	208	

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *-Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
23912	54280	A	24050	182	255	DAESRRSWQPLPEPLELRP\RLS D
23913	54281	Α	24051	146	456	LKNHKRSEKPCPTLTDDIPPQK KCKWPVLALSADITL*KSFSWL ILAQKAPPLSTL*PPLLPAREQM SFD\LIFLYLPKSYKVAPPLSSFT ESLFGLSPPAPR
23914	54282	С	24052	90	215	
23915	54283	В	24053	1	1059	
23916	54284	c	24054	129	215	
23917	54285	С	24055	19	347	
23918	54286	C	24056	228	260	
23919	54287	В	24057	406	602	
23920	54288	C	24058	7	375	
23921	54289	A	24059	322	400	FLLSRVPSV*DPTENQTVQLTW EWLP
23922	54290	А	24060	818	1078	ILDVFSYTMKIEPSPLRSYPISPH YNL*QLPP*LDP*ESGYKTPLSA LLLTFLLYISSFASHKVSSSSVD PLPTCAYLPIGQAHVH
23923	54291	С	24061	1	411	
23924	54292	Α	24062	1	861	
23925	54293	Α	24063	425	484	
23926	54294	A	24064	1844	2097	HSAGTKLRLVNFLLTLRRLVNS VMRLKKSFMMVQRMVSIPKV WSISLGFLLSFVPSVWDPT/ENR TVQLT*HPLPEPLELGPRLSD
23927	54295	Λ	24065	729	807	LFSCSCPPKPPRP*LTAEKGGLC IFL
23928	54296	В	24066	1	1317	
23929	54297	В	24067	509	810	
23930	54298	А	24068		385	MDLPSTVRVTRSSASVMVQGG SEAIRQRQSSAAKPRRSQKESKS LGPEPQGLWEWLPEIYFSQFRR LGNPKSRHQQIQCLMGACFVV TERLSFQDSLDLAEPRHSYPLQ VKRLDLPTVLETETDLEKLDPF TTIPSPVETLQCLFVSLRIKSRVP *FWISQPPEL*EINFWQPLPEPLE LWPKAL
23931	54299	В	24069	1	774	
23932	54300	Λ	24070	350	640	SLQGCLSDNSPTFQRCQTTQGR LPWSFTLSGKSHFSGGVARACY TCQKSGHWA\RNASSPGFLLSR VPSVSDPTENRTVQLTWQPLPE PLELWPKAL
23933	54301	В	24071	85	331	
23934	54302	В	24072	1	1844	
23935	54303	В	24073	1	1169	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
23936	54304	A	24074	63	220	NLTGNIWNREWYSHFDFDLLD PVGFSGSIAKMPKRKLFKCHSV *GHWRSKSE
23937	54305	В	24075	497	898	-
23938	54306	۸	24076	3	678	GGGFSDRDRTIALQPGQQIWM WGIGKQVGGCGWLRGKNRVV GAAKDQSGTGEKFYFIPKAMG AMEGSGQRRTLPDPGAIRNPS GGYSATDCEGSRPAPGEINSHA SHTKPVWWFFTQTRMKFGAVT RIGGLPWEVNPLSSCSLLRE/KS PPTTSGPQSORPKHLT-KFKS A/CYTCRKSGHWAKECPLPGFL LSRVPSVWDPTEN/KTVQLTWQ PLPEPTELLWFRLSD
23939	54307	В	24077	522	873	
23940	54308	В	24078	329	530	
23941	54309	C	24079	147	439	
23942	54310	A	24080	3	3889	
23943	54311	A	24081	2	394	
23944	54312	Α	24082	1	197	
23945	54313	A	24083	1 .	205	CLRAGPHALYLLC/NIPEFTFL\S RADF\ARSQLVDLLTDP\FQ\QEL EELLQVGAVTLILRILPCLPCLP
23946	54314	A	24084	134	182	SCFIPSSQPVAGTERSRFLGPQY FQTPHNPSGPGLGNQLMR*AF
23947	54315	A	24085	257	481	
23948	54316	A	24086	415	630	
23949	54317	Α	24087	123	340	
23950	54318	В	24088	49	270	
23951	54319	A	24089		900	RSRVRREVVALRRGLACIPSLG VWRFPSVGVLAVSDRWCLAVS DRWCLARATLSCFGCAMPMYQ TRRQENDLRTASIAV*RRKQON DH*KQRRWONIORKGPKRYTVI AGNSQSHQPMIFSMLRKLPKVT CRDVLPEIRAICIEEIGCWMQSY STSFLTDSYLKYIGWTLHDKHR EVRVKCVKALKGLYGNRDLTA RLELFTGRFKDWMVSMIVDRE YSVAVEAVRLLILILKNMEGVL MDVDCESVYPIV*ASN*GLASA VGEFLYWKLFYPECEIRTMGGR EGOROSPGAGATFFOLLL

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	1	l		sequence		
		<u> </u>				
23952	54320	A	24090	1	812	MARTPGAELAVAKYPKKGSQA
		ŀ				VHRHSRKQSEPPANDIFNAAKA
	1					AKSDMQHREVRVKCVKALKG
		l				LYGNRDLTARLELFTGRFKDW
						M\VSMIMDREYSVAVEAVRLLI
						LILKNMEGVLMDVDCESVYPIV
		1		l		LFYPECEIRTMGGREOROSPGA
İ		1		l		QRTFFQLLLSFFVESKLHDHAA
		1				YLVDNLWDCAGTQLKDWEGL
1		1				TSLLLEKDQSTCHMEPGPGTFH
1		1				LLGESPGISAIVQGAIEKSGERSS
		1				PPPKRIGSMKEGPMVLPLALDT
						OCWFPVEVPLD
23953	54321	A	24091	152	1209	MPLKNVQNITHFPLEKELALPL
23,33	34321	ľ`	24071	1.52	.20)	LIFSPLHWCVSGYLSLPLPNPKP
						HFSNPLLLQVILPALTLVYFSIL
				ŀ		WTLT\HISKSDASPGECGS*MG*
						WEQRNLCPSILPLQSPLCPTEAA
		ŀ				VEFEGTRMVAFCELC\QSCLSD
1						VDTEIP/ESR*VLAOSGRKKAEY
						PCPPFPRRN*QCHYLPWETFCPF
1						KHPFFSPLQAFVLLSDLLLIFSP
		1				QMIVGGRDFLRPLVFFPE\ATLQ
ŀ						\SELASFLMDHVFIQPGRPGQW
				1		CSDSIPGAQ*GLGLGCSSKI*GR
	Ì	1				KLKSVIPFCLPCEPGDSQEDHLQ
						IERLHQRRRLLAGFCKLLLYGV
						LEMDAASDVFKHYNKVHQGPS
		<u> </u>				LGFTKPLVMPRFSAATV
23954	54322	A	24092	37	625	SLMLLTPFHFPPEECPPPIPTAH
İ						HLLPFLLSTWELFLQQLQEVVV
l		1		1		KHAEPAVLEAGAHAL\YLLCNP
1		1				EFTFFSRADFARSQL/V*DLLTD\
						RFQQ\ELEELLQGRELGAGTMG
ì		1		1		DTPNKGWGGGCKTGGMWDL
1		l				MSSGGKCYCGIFFSFSILPSQIED
						EVYNLGSHSETPLCLLQVSGFP
						PLPHPVFTGVCGVGSLCSFPV
23955	54323	Α	24093	2	198	PNDRSRVTREMSAPSRSLGRFL
		l				SVGVWHAPPSLALVPPCRCTRL
1	1	1		1		KTYHGRGAPLRVELPTCMYRL
						SNVQGRSGGPAPGVGHLQSLV
1		1	1	I		DEWLDSYKQDQ\DAGFL\ELVN
1		1	1	1		FFI*S\CGM*RNSGDYPLTAPGL
1		ı	l	1		SWKKFQGSFCEFVGTLVCRCQ
1		1	1			YILLHDDFPMDNLISLLTGFSDS
1		1	l			QVCAF\RHTSTLAAMKLMTSLV
1		1	l			RVALOLSLHODINORQYEAER
		1				NKGPGQRAPERLESLLEKHKEL
1		ı	1	1		H*YQAKNLSRARCASTCGASH
23956	54324	В	24094	211	3841	Q. Markold McAdire GASII
23730	J4344	10	24074	1-11	50	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23957	54325	Α	24095	1	1798	MSLLAFKIILVGFEGSKARVQT
		1		l		DLFYRSLVLFLLLQICFPLSCFG
						NWKGEEGGMQRWVLGKNEKI
						ASPHRLPAPTARFAVSLQLNLRI
		ł				LGAFQFGPVLFVLWRPLDRDN
	İ	ı				ASPQSEFLNSCGITAYLVGKRE
		1				GGVERGGGGRSSERCRTRLSFS
		1				PRHSHVFSTSRFCKILKKSPGAN
		1	İ	1		GGRCRARGGGLHGARAANPPT
		1				PIAEPARLPACLSRAEPEPQAAE
		1				PAPPPVGVVRSRPLSHRRHPVV
		i		ł		CPAAPIRLINKRLAHSARPAGIC
						CVSRSGLSALAAAGTASMFQLP
		l				ILNFSPQQVAGVCETLEESGDV
				•		ERLGRFLWSLPVAPAACEALN
						KNESVLRARAIVAFHGGNYREL
		l				YHILENHKFTKESHAKLQALW
			l			LEAHYQEAEKLRGRPLGPVDK
		l				YRVRKKFPLPRTIWDGEQKTHO
						FKERTRHLLREWYLQDPYPNPS
		l				KKRELAQATGLTPTQVGNWFK
		1				NRRQRDRAAAAKNRSQTSPWA
		i				EAFRFPECGSRLAKHFAEEPGLI
		ĺ				PQAQMLLTILFAFRSGRPDPTRS REHLLRRKPOADAPLEM\LOOO
		ŀ				VLSOGSGRALRAEGDGTPEVL
		l				GVATSPAASLSSKAATSAISITS
23958	54326	A	24096	3	1034	GLHGARAANPPTPIAEPARLPA
23936	34320	n	24070	ľ	1034	CLSRAEPEPQAAEPAPPPVGVV
		l				RSRPLSHRRHPVVCPAAPIRLIN
		l				KRLAHSARPAGICCVSRSGLSA
						LAAAGTASMFQLPILNFSPQQV
		l				AGVCETLEESGDVERLGRFLWS
		1				LPVAPAACEALNKNESVLRAR
		1	ļ			AIVAFHGGNYRELYHILENHKF
		l				TKESHAKLQALWLEAHYQEAE
		l	1	l		KLRGRPLGPVDKYRVRKKFPLP
		l	1			RTIWDGEQKTHCFKERTRHLLR
		I	1	l		EWYLQDPYPNPSKKRELAQAT
		l	1	l		GLTPTQVGNWFKNRRQRDRAA
		l				AAK\NRLQQQVLSQGSGRALR
		1				AEGDGTPEVLGVATSPAASLSS
		I				KAATSAISITSSDSECDI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			1	sequence		
23959	54327	A	24097	3	11024	GTRVGVAWRSVRLLLGPGAGL
23939	34327	Α.	24097	3	1024	RGGVVLKVSPSPCRGGRRVPVR
1		l				AEGARGRGPINKVPGASSGAA
1		l				VGAARLTAPLLAKAMASKLLR
						AVIL\GPPGLGKGAPVCQR1AQN
1						FGLQ\HLS\SGHFLRENIKA\STE
		l				VGEMAKQVYREKVFLVPDHVI
į.						TRLMMSELENRRGQHWLL\DG
					1	FPRTFGQAEALDK\ICEVDLVIR
		1				LNISFETLKDGLN\RRWIHP\PSG
						\RVYNL\DFNPPHVHGIDDVTGE
				i		PLVQQEDDKP\EAVAA\RLRQY
ł		l				KDVAKPVIELYKS\RGVLHQFS
1						GTET\NKIWPYVYTLFSNKITPI
				l		QSKESILTLAQWEEPGRCGSFIP
						IVWCSNWCCCPNLEA
23960	54328	Α	24098	2	178	
23961	54329	Α	24099	96	340	FGALTRIG\DLPWEINPLSSCSLL
		l				PEKDPPMTSGPQTDQPKEHLTN
						FKSAARRSSRTSSHRNLLHVLEI
1		1		i		WQLDQGMPAARDS
23962	54330	Α	24100	3	155	
23963	54331	Α	24101	1	162	
23964	54332	Α	24102	1	1032	
23965	54333	A	24103	167	476	TAMLLTQSLFGGLFTPTHMKFG
			ļ			AVTQIG\DLPWEINPLSSCSLLR
				l		EKDPPTTSGPQTHQPKEHLTNF
						KSDHHGCRASGNSHSGRMLLIF
						SPSRGLYACSCVKT
23966	54334	A	24104	371	772	
23967	54335	Α	24105	146	312	QKRSIPNSSPIP*SHSLLANCTK/
		l				DHHFLNGSKSFQPQLLSCKFS*
						GQHLFQGALCT
23968	54336	A	24106	76	441	
23969	54337	A	24107	1	1284	
23970	54338	Α	24108	150	371	EESRAAGIPWPSGQISGTCSKL\
						CGRRFWRNAWPLRFRHLEVLV
				i		CWRCGCGLSHSGGKELQLFSII
						VHLEGEVN
23971	54339	Α	24109	280	351	
23972	54340	A	24110	1	483	
23973	54341	A	24111	129	363	FGVVTQIG\DLPWEINPLSSCSL
			l	1		LCEKDPPTTSGPTKKHLTNFKS
				·		SKRPLFTLFSNLPHYPSTSFSFQS
						WRYTSISPFS
23974	54342	Α	24112	2	111	
23975	54343	Α	24113	3	808	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (N=Unknown,Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion) TALLLTOILFGGLFIWTRVKFG
23710	34344	,				ALTWIGDLPWEINPLSSCSLLH EKDPPTTSGPQTDQPKKHLTNF KSACFERIKACYHSPATAWPFK AYKLSLQFPHFTCPKTR
23977	54345	Α	24115	ı	753	
23978	54346	Α	24116	1	513	
23979	54347	Α	24117	625	701	IHVAHKLAVVMPGCKRGCLSL FYFQLIWSHKGWVRSGEASHR EMAGLLLLCSS*PPSPCETLPRS EPSLCDSRSAESRTKTSNLSCIP ASLQQVCGPREFIWSGNHCHPF LPITRCYGCLCYHYCMENSIL
23980	54348	A	24118	458	605	VVEEGIEDKRVYRLGTTGWIGK TIWLVRRRS*TNL*DLSGFWTG KMGEL
23981	54349	A	24119	566	769	IKEKGGLFSFSASNLWKQVTSS FLCSPLPLSLLKSSKLLANRDKY RMCGPVPANGNRTQQ*SKHVR
23982	54350	Α	24120	345	664	GEIAASQEPAGGFSGSSAQPCPL PWTGPPTTQAPGH/CNSAPSSRD NASSNGGRRQRVPTVPGGGLIL QREHILLPLGPALRAAGKEPAT AATIPESPGGDSDDSRT

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SEO ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	i	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		l	l	sequence		
		<u></u>			L	
23983	5435I	A	24121	2	2204	IDMIFTPGPPSTPKHKKSQKGSA
						FTFPSQQSPRNEPYVARPSTSEI
				l		EDQSMMGKFVKVERQVQDMG
1						KKLDFLVDMHMQHMERLQVQ
						VTEYYPTKGTSSPAEAEKKEDN
1						RYSDLKTIICNYSETGPPEPPYSF
1				-		HOVTIDKVSPYGFFAHDPVNLP
			İ			RGGPSSGKVQATPPSSATTYVE
1				l		RPTVLPILTLLDSRVSCHSQADL
1				i		QGPYSDRISPRQRRSITRDSDTP
	1			ł	•	LSLMSVNHEELERSPSGFSISOD
					İ	RDDYVFGPNGGSSWMREKRYL
					1	AEGETDTDTDPFTPSGSMPLSST
						GDGISDSTVPLFLSSEILOKOVG
						QSITSMLGFLSRGPSMKLCMGL
						ACVLSLWNTVSGIKGEAKKEK
		1				GMTFLPTTDSKKFFSLLSVTSYS
1		1	1			SFAFHKFSVAVYNISNLKTVDP
						AKFPTRYCYCLNNRTNDLSDFT
						ALLVDIIGNSTSYLTEIFKSTSIL
						SVNQSNESDCIFICVMTGKSGR
						NLSDFWEIEEKYPIINYTFTSGL
						SGVLALLLTQSLFGGLFTRTRM
		1			1	KFGAVTRIG\DLPWEINP/LSSCS
		l		l		LLHEKDPPTTSGPQTDQPKKHL
					1	TNFKSAARPTFLGQGQVPLNPF
						SFTLSGKSCFPRRQEPPNRLFPH
1			1			PNLLSLCPNPLFLCPNPFSTFLE
					:	GKNHHPHPFSVSQVFSLGLPPS
		ł				LWEHSPMAHRLRVOPROOPPDI
		1				HKQVISLQSGHSLLVKSQSWSQ
		l				DPIKFQLPHHHKNTASVCINPSV
23984	54352	A	24122	2102	2529	ETTMSFIFSLLYPFRMHLSFLIVS
23704	34352	ľ	24122	2102	2027	PALRSISRRALAAFPRGRSLDLQ
		1		1	1	PAMPEPPTPS/CGLLCVPS\PPMS
		1	1	l	1	
			I			AAPCSKVPQSHLFAFMSCNTAK VCSFTAEASETTSPPGGTNNSR
						RATLRAVTLTRRSATLTRRSAA
		L				SHRSSTV
23985	54353	Α	24123	299	567	TAMLLTQSLFGGLFTQTRMKF
		l		l		GAVTQI\ADLPWEINPLSSCSLL
1		1		1	1	HEKDPPTSSGPQTDQPKEHLTN
		1				FKSGETKEMHFIRGPKTPAPVM
23986	54354	Α	24124	684	822	KVKRQSYSQNTFYYPICSRH*IK
1						LOKLNSGPOTPOODLINLAIKV
		-				

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide scquence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, *=possible nucleotide insertion)
23987	54355	A	24125	1163	1826	TVLLLTQSLFGGLFWTRMKFG AVTRINGBLYPWINPLSSCSLL HEKDPPTTSGPQTNQFKELLTN FKSYPTTSVAESFQSSFSTDPSD PSSPPQAARPQAKFGPNSSSASA PRPYNPFITSPHTWSGLOFHSA TSPPPPAQOFLKKVAQVKGT KWLKTDAARSPQKPTGTTTDAL DNSHSGGFIMDRFLLVKSPKGF LRLLYFSKTFISTILINLQE
23988	54356	A	24126	636	922	TATLLTQSLFGGLFTRTRMKFG AVTRIGDLPWEINPLSSCSLLRE KDPPTTSGPQTHQPKEHLTNFK SDKGDMFYPWTQNSSAGHGLG RQPSLGV
23989	54357	Α	24127	728	1013	MKFGAVTRI/RGDLPWEINPLSS CVLC/CREKDPPTTSGPQTDQPK EHLTNFKSGKRPAFYSLLQPPSL SLNLFLLSILAPHFNLSLLLISIPF IFW
23990	54358	В	24128	I	3993	
23991	54359	В	24129	105	420	
23992	54360	В	24130	45	2866	
23993	54361	В	24131	I	1680	
23994	54362	Α	24132	1257	1403	
23995	54363	Α	24133	1	2130	
23996	54364	Α	24134	1	725	
23997	54365	A	24135	2825	4123	RAADGVITLEDNDPCVKDDSSR FPYEDRLDLVLKGTADIPRLTV HRGSEYIISRATFFCYEIKEQSVI NHCYTEIDLKIFRGYLAPALGV HREFYGTEPCRVTAQYNQDM RYWLETPTISAPPIELVEIERLRY QEMPIYASRWHFIERDDTALP RNDTPOMVKLFMDREQVVKDI GYIELKVVEDQRTRAVMIKFRA FVKEGA VIVIRFDNKEVAFAQM SGNLTVPRSTSDYKARDIFFGY REITHISTIGHGVANNHQIRLRL ELAGIVPLNGJNALLLKVYPLS KLCNSPKKCQLESVFRLTQWV VKKPISNRQSGLGGIVLPGLFRV LSPGNLSPIKGTVILPNEVQFGF TGGNLSPIKGTVILPNEVQFGF TGGNLSPIKGTVILPNEVQFGF TGGNLSPIKGTVILPNEVQFGF TGGNLSPIKGTVILPNEVQFGF TGGNLSPIKGTVILPNEVQFGF TGYNSAGLADHTRASSPERA GVYPHHRECAHRASSPERA

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nuelcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /possible nuelcotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
23998	54366	A	24136	137	1199	ISSGGPRELGLKMIIVPGSQLSQ
		1		\		LPPGLDSTEFAAIVGNLLDNAF
						EASLRSDEGNKIVELFLSDEGD
		1				DVVIEVADQGCGVPESLRDKIF
						EQGVSTRADEPGEHGIGLYLIA
		1				SYVTRCGGGITLEDNDPCAPLT
		1				LLIVEDETPLAEMHAEYIRHIPG
		l				FSQILLAGNLAQARMMIERFKP
						GLILLDNYLPDGRGINLLHELV
		1				QAHYPGDVVFTTAASDMETVS
	ľ	1				EAVRCGVFDYLIKPIAYERLGQ
		1				TLTGFP\QRKHMLKSIESATQKQ
						IDEMFNAYARGEPKDELPTGID
İ		l				PLTLNA VRKLFKEPGVQHTAET
	1	l		1		VAQALTISRTTARRYLEYCASR
		1				HLIIAEIVHGKVGRPQRIYHSG
23999	54367	A	24137	2	243	RPGRGLGQRGIWEPAA\LLDSV
						AQQQGAWLPHVADFRWRVDV
						AISTSALARSLQPSVLMQLKLS
						DGSADLEKRCERRLQD
24000	54368	A	24138	1	1483	MKLMETLNQCINAGHEMTKAI
		1				AIAQFNDDSPEARKITRRWRIG
						EAADLVGVSSQAIRDAEKAGR
						LPHPDMEIRGRVEQRVGYTIEQI
1					l	NHMRDVFGTRLRRAEDVFPPVI
		1			i	GVAAHKGGVYKTSVSVHLAQ
ŀ		ı				DLALKGLRVLLVEGNDPQGTA
	1	l				SMYHGWVPDLHIHAEDTLLPF
		l				YLGEKDDVTYAIKPTCWPGLDI
		l		1	ŀ	IPSCLALHRIETELMGKFDEGKL
1		1				PTDPHLMLRLAIETVAHDYDVI
	ł	1		1		VIDSAPNLGIGTINVVCAADVLI
		1		1		VPTPAELFDYTSALQFFDMLRD
		1		l		LLKNVDLKGFEPDSDLRYLQSC
i		1				EWGDTGIFLKWSCIPAEKGLQT
		1				SGAWVEAADIPLLPGSPRRLSP
				1	1	QAGSRGGQGPKHGQQCLKMP
l		1			I	GPRAPGLQGGSNRDP/AP/GPAV
l	1	1			1	ERAPAAALSSTTIWMPMSLCP
					l	WRPVLAACTSMTVRGRSTMYL
					1	PTTTGNAGCTWPKASLATRWL
		1			1	SSPMGRQARSLMLGAQVTLSW
	1	L.	l			DRSMPWRRRGRSSGRNLSTI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*-Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24001	54369	Α	24139	727	1498	GICIWDRPPGLADQRRSHSHGW
					1	WGELATPLPWHH/HLSDSHSG\
		ĺ				RVSFLGAQLPPEVAAMARLLG
l						DLDRSTFRKLL\KFVVSSLQGED
İ						CREAAAAVLGSAPTCRR\AAGC
					1	
						LLAGMHTLLQQALRLPPTSLK
					l	PDTFRDPAPGALHPPRPGRGL\G
		1				QRGIWE\PSGPSFDSVAQQARV
		1			'	AWLPHVA\DFR\WRVDVTIST\S
	l				i	AL\ARSLQ\PSVLM\QLKAFQNG
		1			İ	SSIPAFEVPPQPKF\QEL\RYSVA\
		1				LV\LKEMADLEKK\CE\RNLQD
24002	54370	Α	24140	1	312	
24003	54371	Α	24141	2	524	
24004	54372	A	24142	144	494	AGSCPFAAGPGPLGSGRCFVLIR
	ŀ					FILI\ONRAGKTRLAK W\YMOFD
		ŀ	·			DDEKOKLI\EEVHAVVTVRDAK
			ŀ			HTNFVGVSGTFKIILPPLCWPLL
	l		ĺ			LASVWMFNDNKPGLYLGGPFH
			l			NFRGRS
		ļ			700	NEKGKS
24005	54373	Α	24143	1	428	
24006	54374	A	24144	108	422	
24007	54375	A	24145	3	387	
24008	54376	A	24146	3	337	TRLMLLDRSGQGKVYGLVGR*
		ł				RFQQMDVLEGLNLLITI*RNRN
i						KLRVYYLTWLRIKIMDNDPKLE
			ŀ			KKHDWTTAGYMEGCGHYRVG
İ		1				KYERIKFLVIALKSSAEVYAWA
	ļ		ŀ			PKPY
24009	54377	Α	24147	1	340	ERTRMNKQHNSTLAKSRPGNT
	1	1	İ		1	GPEPLMPQASPGPPGPLSQTPP
	1		i			MHRPVEPOEGPHKSLVAHRVP
	l		l			LKPYAAPVPRS\QSLQDQPTRN
	1				1	LAAFPASHDPDPAIPAPTATPSA
1	1	ĺ				RGAV
24010	54378	A	24148	335	869	ATPTLILSOKSHFWPOPLSSVLK
24010	34378	l ^A	24148	1333	009	
	1	1			l	LTSGPEISLQSSPSPICRMKKRR
	1	1			Į.	LSPLASGLES*VKSGEWCEVLS
1	1	1	4	l	I	RAPLFP/HNPSQAAP/EWEMQG
1	1	1			l	TRPSIPHPLGLAGLPTQLQQGRL
1	1	1			l	WGYRNPASPFRTRVAARDCAG
1	1	1	1	1		KQVRTGPDARLEGSRPGVRAG
1	1	1		1		LGWGRGYSERVQTCAKSSVSPS
24011	54379	В	24149	158	5640	
24012	54380	A	24150	ī	1854	
<u></u>						

SEO ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	
1	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24013	5438I	Ā	24151	1	1510	MPCGKQGNLQVPGSKVLPGLG
12.013		l''		ľ		EGCKEMWLRKVIYGGEVWGK
1		l		l		SPEPEFPSLVNLCQSWKINNLM
1		l	ŀ			STVHSDEAGMLSYFLFEELMRC
						DKDSMPDGNLSEEEKLFLSYFP
		1				LHKFELEONIKELNTLADOVDT
1		l	ŀ		i	THELLTKTSLVASSSGAVSGVM
İ					l	NILGLALAPVTAGGSLMLSATG
1						TGLGAAAAITNIVTNVLENRSN
1					l	SAARDKASRLGPLTTSHEAFGG
		1				
						INWSEIEAAGFCVNKWVKAIQG
		1				IKDLHAYQMAKSNSGFMAMV
			1		l	KNFVAKRHIPFWTARGVQRAF
i						EGTTLAMTNGAWVMGAAGAG
						FLLMKDMSSFLQSWKHLE\DG
					1	ARTETAEELRALAKE\LEQELD
					ļ	RLTQHHRHLPQKASQTCSSSRG
						RAVRGSRVVKPEGCLATSQPH
1						QHQQERGDRPREDTDNEKRGG
}		ŀ			l	EKWSDFGSVLKGEPTGFVHGL
1		l				DLWCEGKKSQDNGGLWPEQL
						AGWSCWHLKWTGGSRSGLPVP
						VLDAACTGTLSPDLTSWRYHPS
		_				SIPLPKQPLQHGF
24014	54382	A	24152	3	515	VGTQAGAAPGPVQFLSSAWVA
1						L\HETELPPLTAREAKQIRRE\IS
1						R\SSKWVDMLGDWEKYKSSRK
		1		1		LIDRAYKGMPMNIRGPMWSVL
i						LNTEEMKLKNPGR\YQVRSARA
i		1				QPTGQAVSGAQVSSWRERQDH
1						PGELGVKIMKEKGKRSSEHIQQ
						MDLDVSGTLRRHIFFRDRYGT
24015	54383	A	24153	1	1317	
24016	54384	Α	24154	1640	4841	TITGSSPLAGGHHSTCRGRMDM
			1			VENADSLQAQERKDILMKYDK
		1				GHRAGLPEDKGPEPVGINSSIDR
			1			FGILHETELPPVTAREAKKIRRE
		1	1			MTRTSKWMEMLGEWETYKHS
						SKLIDRVYKGIPMNIRGPVWSV
	1		1		1	LLNIQEIKLKNPGRYQIMKERG
		1	1	1	1	KRSSEHIHHIDLDVRTTLRNHV
				1		FFRDRYGAKQRELFYILLAYSE
1					1	YNPEVGYCRDLSHITALFLLYL
1						PEEDAFWALVQLLASERHSLPG
1			1		1	FHSPNGGTVQGLQD
24017	54385	A	24155	1640	4056	, = ,=
	12					·

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
24018	54386	Α	24156	479	3902	GQVDEDV\$VKMDTVPGVNLSC
				l		ILNEMRDQDKKLVEKSYKDAK
						GCFFSMGEPSLIDPCLEVCYGT
						QLAQLQGLIRSMEQQLCELCCD
			1			AEHQDHEHQVLLDVKTRLEQEI
						ATYSRLLEVEDAQQGPIVMRA
						MTSECHWCLAPIRSPDCVCVLR
						QLHTRSGIHKRFCGLKGFLPER
						HNPGQLIHQNQSGCCSRMYPFD
						RKLGISVALNIPGGRSYRDLWC
						LTAVHIHIPGIDHHRLFTLGRM
						DMVENADSLQAQERKD
24019	54387	A	24157	780	983	QHQSLIGHKYQRSHRSL*AEPE*
						LECHQPDLAEPADGPPLTTRYG
						IVNAEMGLEQLLPKGPSLKEAK
24020	54388	Α	24158	773	2016	DWPTLRENRKEPTLKYWGLVP
		l				LISGMPTWFFFFLSACGNLIPSG
		l				GIRVPMENML/RVFRAVEEHCP
			ĺ			WFPEKGTLSVEQWDSVGAKFQ
			ŀ			ELVPTGNYVPITVWGDWALLL
						QFSESGDPLPLPQLSSPTGPSLS
						DQPLPSPTPPPPDDVENSISNSG
			l			DFGLTLPPGDLILFPEEPLLAAS
						AAPNRTALGHIYANSSLFKPLQ
		ŀ				HLPLESANGSGAKLQFTYNSAG
		1				PSPSSAAPRPPVVSVPQPVTLPS
						TQAASLYPSSHMDTSNHQCTSA
						SSAPPMPLSHTLIPSDETTCNVF
						NYPLKKPTGRELREAKNKGSQP
	1	ĺ				KVHEKTNPGNIRCISTEERPHEN
	l	l				KGFNHLLEWLAELKTTIHLLDC
	ĺ					WFIAKDVNREQPDGRDAGDKR
						LCVKFLREIAANTYKTEVGAAK
						VMEILLHSHGATTTIAN
24021	54389	Α	24159	365	675	
24022	54390	A	24160	I	1191	

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SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nuclcotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	İ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
24023	54391	IA	24161	1363	1434	QQGVCSGEKGGHWAPROVGV
						YLLPGRVGCVSSRVSPSFPGDG
1		1				LDSGLARRGSAVSALASGLVEE
		1		1		PMLGPPFHPTPRFKAVSAKSKE
						DLVSQGFTEFTIEDFHNTFMDLI
		1				EQVEKOTSVADMLASFNDOST
1		1				SDYLVVYLRLLTSG\YLQRESQ
		1				VLRALHRRVDGTVORSSAKOE
		1			l	VEPMCKESDHHSHHCAGPGPS
		1				ACPSRVGVHGTAARAAPPIRTS
		1				F/H*GVPSPKVYLLYRPG\HYDI
		i				LYKLGLGSSPLGCPGCPLLARA
		i				LGHCYRGFSVVVNGPISPPSSCH
		l				MTPPPCFIKGGAGGEPCVRVPA
		l				LLPAWLLCLLPPPPRWVPLLFT
		l				YLLLSFPNRSRFEGPGLLEAPPA
		1				SLGSASFPS
24024	54392	Ā	24162	2	888	VFRSRHOGCLKMAAEEPOOON
24024	34332	^	24102	-	800	OYPLGSDSEGVNCLAYDEAIM
						AQQDRIQQEIAVQNPLVSERLE
						LSVLYKEYAEDDNIYOOKIO\D
						LHKKYFVHPQDQGLTGTCFYS
		1	İ			GFRDSPTWEGTCWIDSKELORF
ŀ		1	İ			QGLVS/SQRSKGKTLVSQGLSL
		l				EFHNFEGFSTTTFI/VDLIEQVEK
		1				ADLCRRPCLASFNDQSTSD\YL
						VVYLRLLT/SRAYLORESKFFEH
		1				FI\EGGRTVKE\FCQQE\VEPMW
		1				QGEATTSTSLALAQALSVSIQV
l		1			ļ	EYMDRGEGGTTNPHIFPEGSEP
		1				KVYLLYRPGHYDILYK
24025	54393	A	24163	3	819	EGATMAAPPOPVTHLIFDMDG
i		1			1 '	LLLDTERLYSVVFQDI\CHRYD
1		l				KKYSWDVKSLVMGKKALEAA
		l		l		QIIIDVLQLPMSKEELVEESQTK
		1		l	ŀ	LKEVFPTAALMPGAEKLIIHLR
						KHGIPFALATSSGSASFDMKTS
		1				RHKEFFSLFSHIVLGDDPEVQH
		1				GKPDPDIFLACAKRFSPPPAME
			l			KCLVFEDAPNGVEAALAAGMQ
1		1	l	l	l	VVMVPDGNLSRDLTTKATLVL
1		1	l	l		NSLQDFQPELFGLPSYELEGRA
				1		SVFRPQPTLMVHTAGGKGKEIS
						NSSIPTCAVILAS
24026	54394	В	24164	115	594	
24027	54395	A	24165	4	160	SIIFPAIPVSLK*IKPNCPEPPPDH
				1		LLQAAWGCVSLGQGLSYSALN
		1	l	I		TLLYFT

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24028	54396	A	24166	88	2280	VPLYDRMGGETGLLIPSPERSL HAATDAHSCKNRHSLTGKTDS LWQQDTKLQTRPKATQGKGES GGCGPHRPLLLLRTKSTPESKS AHDPPVAPAQAGASGCGRGAA QKGFGAKAVDGGGRDRTPG VPAAGPPTSSGRGDRGPSSSRR
						TRSQTPESPTGRPGSRQFQAAPP PSHHSRPAHAPFPGFQMYRGPL YSSSPAPATPNTSAYESPRFCST SYISSSPRFCPTSYICSSPRFCSTS YICSSPRFCSTSY
24029	54397	A	24167	2	352	WRTLIVLGMSCLGSYPK*MHLT HLIHCS*EARSLVTLHIPLTICG EPRNIIKLVGCP*IQWKK**KKI MNSRILSPSFRSRY*TSNLLGLP *LRGLSPVERELKLWTNRHKLL
24030	54398	Α	24168	2	252	NKSSVFATSCSLDPISIMSLM*W TSVISCRSKK*SRSLRIRL*HKA KELIYPLPVRESMWGYCQLLS MSMPIMHAGTGEMTTG
24031	54399	A	24169	1505	1680	RTSSLVTL*TLHLEPAGT*VLTP GSTTGITVSGTVLFLILLSCFLSS TPFRLKLACLS
24032	54400	Α	24170	681	932	
24033	54401	А	24171	1740	1950	HHHHLLQMSWSAGMQAWHW PSFYNVVTGKTLA/LPNLIALQH IPLSPAGVIAKRPAPIALPNSCA ALLSSQ
24034	54402	A	24172		652	GKLITEARTRCCMQLTSDRINGH TALNHTTCSHICVSGMAGRL GPITVLSGD\GPKGMRALTMW HTLARPTGDLEAPRHGPRRIT ATOPPDHAPPATSRINRAAHPHN HTPPHPQTAHPFATHSHLTYR TPRPTPATRTPLHPHSHQVPY Q\text{QPTQRSPYLHPPPAPPHISLHPS} TTRATVNAPDTRFTIYARESVPF GRILINLGRQRGMELANILQVH GP
24035	54403	A	24173	267	400	RFLHCGNLAGLLVGFLSFKLGE YEGLGPLAALTAGKGLFFLPLE SRLRLHRRPKPRSPASFG*VSRH THAIHFN
24036	54404	Α	24174	422	452	
24037	54405	В	24175	131 .	1047	
24038	54406	Α	24176	2	199	LGLR/CARLEARARPPWATVRE KLWLFGHPTHMRCSWRGLFQD DLGLQKPLSGGSSSAYRGCENL NS

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod		location of first		*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24039	54407	A	24177	440	1746	RNLOCVFOGRIKEGIPVLALS/L
		1			1	RPSP/RVLHTATKQKNSCQHLE
						EDV D\QTKWRTDTPSMDKILME
		l		1	1	EVKLEEQLKEAVEEDKQALAD
		1		1		TEGSEQSSQKLVEEGNMYSIQG
		l		1		FCKDSLEVADVLEKATQCVPER
		1			1	EIKDNNPHLKNLCETLTMSEEL
		l				SQMVDCCRLPQDVSVLGVALH
						WVTPQMLCVPVNEDIPEVSDT
		l				VLKAITAPPRVPIGRTSGDVTGT
		i				GVRRRRVSSAMSQHRAFRDVT
	}					VTVQCLELGCFLLRVELASQFL
	1	1				KRPRSPESNEKVPEIEVTVEASA
	1	1				KMSHPTLYDATRSPWGMFQSQ
		1				SVLLSCAHHSLLRGATLDRCGS
						IDESEEWYGMDTFECTGVGVSI
	ĺ					RRATEPERSSGSAAALDSHRSM
						HPIVNHACEGSTLRCPYENLMP
		l		l		DALSLSPILSSRSGDASPQAEILF
		l	1	l		VVGPFRMRRCLTVDPPGLDVT
24040	54408	Α	24178	391	656	

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SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24041	54409	A	24179	1	2150	MEDSTSPKQEKENQEELGETRR PWEGKTAASPQYSEPESSEPLE AKQGPETGRQSRSSRPWSPQSR
					1	AKTPLGGPAGPETSSPAPVSPRE
		ł				PSSSPSPLAPARQDLAAPPQSDR
		ļ				TTSVIPEAGTPYPDPLEQSSDKR
		1				ESTPHHTSQSEGNTFQQSQQPK
l		l				PHLCGRRDVSYNNAKOKELRF
		l				DVFQEEDSNSDYDLQQPAPGGS
		1		1		EVAPSMLEITIQNAKAYLLKTSS
						NSGFNLYDHLSNMLTKILNERP
						ENAVDIFENISQDVKMAHFSKK
		1				FDALQNENELLPTYEIAEKQKA
		l				LFLQGHLEGVDQELEDEIAENA
ļ		l	l		Į.	LPNVMESAFYFEQAGVGLGTD
1		l				ETYRIFLALKQLTDTHPIQRCRF
		1				WGKILGLEMNYIVAEVEFREGE
1		1				DEEEVEEEDVAEERDNGESEAH
				-		EDEEDELPKSFYKAPQAIPKEES
		1				RTGANKYVYFVCNEPGRPWVK
		1				LPPVIPAQIVIARKIKKFFTGRLD
		l				APIISYPPFPGNESNYLRAQIARI
		1		l		SAGTHVSPLGFYQFGEEEGEEE
		1		l		EEAEGGRNSFEENPDFEGIQVID
		1				LVESLSNWVHHVQHILSQGRC
						NWFNSIQKNEEEEEEEDEEKDD
1			ļ.			SDYIEQEVGLPLLTPISEDLEIQN
l		1	l			IPPWTTRLSSNLIPQYAIAVLQS
		1				NLWPGAYAFSNGKKFENFYIG
		l				WGHKYSPDNYTPPVP\PPVYHE
						YPSVPEMAEMDDPSVEEEQAF
						RAAQEAVLLAAENEESEEDEDE
24042	54410	В	24180	1	489	
24043	54411	Α	24181	39	423	RSPTPDAASRSQCSSRRRRGVS
						KEVFVSMARTKADWPANRTG
						G*STPRKPTGPTKSPLRKSAP/A
		l			1	PTGGGERNPHRLQGLVTVAAP*
						NLDGYQKVHSNFLIRKLSLPSV
						WVREIASGLLKQNLRFQERQ
24044	54412	Α	24182	3	384	
24045	54413	A	24183	1	434	MLQNVTPARAEWQRKTALQG
		l			1	VEDPVPARPPISSPEDLLRSGEC
1						STAPTQRQVPQPVHE/SHRPTQ
1		1	I	I	1	DSPGARRKPTSERGHTEQRAAP
1		1		I	i	DREARPATRGWGLRPGGQWLG
			1			GSEHAAAAPGPSDHPRGRCRH
		L_	L	L	L	HTILQRPRLLSGKTES

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24046	54414	A	24184	15	286	LPSRGAWLGTCSPPCLSLPPPP WAPVRPEPPR*APPPAPRRPVPS TTQGSLGPHC*PLE*HSPRRSA ASLLKPARPRAHQEERTTPDAP P*EL*HSP*RSAASLLKPARPRA HQEERTTPDAPP
24047	54415	A	24185	2	214	CLSLPPPPWDPVRPEPPQ*VPSP APQHPVPSTTQGLRSVGARRGT GRQLHLQSRCGIHWVKPAGLL SLGL
24048	54416	A	24186	308	311	LPSHGAGLGTCSPPCLSLPAAP WAP/RVARASPTSAAPTPQHPV PSTTQGPLGSPSTLRKL*SFALC NKSCCCSLFGSTLPL*AVTLTA KVCSFTPKA\ARP
24049	54417	Α	24187	1	669	IMMPEGSGISLSGCDLGTALQLL SQAIVECLKLFQMLNRSTILGSG GQQPSSHSSTKLRPDSGAQLAS PSGSCTGAAGGAACQSRTVCSH SSALGWSMGLGAVEGGVVSLV GEAQAAQEPMEGVGSGMAG CRSRALPHGKAAKARRAIEHST GGLALLGDPVBSPQFLARVLSP SLPRASRAGWPAPSACPPSPHP PGTPAGPQAPHAAPVPARASPS TPPCKL
24050	54418	A	24188	157	395	RCSSCQREMTPYGK*NVQKRL KTTKNGSM*LKSHILSTPEQ\PA SPTSS/PPLLQPAAPAPPLMPRPS PPPGISVA*YPGM
24051	54419	Α	24189	2	287	CLSLPPPPWAPVRPKPPR*ATPP APRRPVPSTTQGLRSAGARGT GRQLHLQPLCGIHWVKPAGLLS LVGTWRTFMSSSGIVNTPISTLC LAQGL
24052	54420	A	24190	500	724	
24053	54421	В	24191	1	2154	
24054	54422	А	24192	688	1215	SRLAPALLSPWVVDGTGRCGA VGGAHWGGLGRTGAHGGGR LRHGGLRVPSPAPREGS*GPPR YRAQRWWAGT/DWRTQYTLRS CWFGTLPSK*FEMTPYG*NV QKRLKTTKNGSM*LKSHILSTP EQWASPTSS/PPLLQPAAPAPPL MPRPSPPPGISVA*YPGMPWTC PFYWQ

SEQID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	eodon for last amino aeid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	eodon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
		<u> </u>		1000	<u> </u>	LIVER THE PERSON NAMED AND ADDRESS OF THE PERSON NAMED AND ADD
24055	54423	A	24193	308	1131	VLQLLRRRTWSCSLRAWSWDL
		ı			i	IVPPGGLVVSLASGVKLQTFAV
						SVTALKEARLDLFVPPGGLVVS
						LASGVKLQTFAVSVTAHKSSV
		1				DPKNSGAQLASPSGSRTETAGG
	i	l				AACQSRAVRSHSSALGWSMGL
		i				GAIEQGAALVRGGSGRTG\PME
						WVGGSGMAGCRSRALPRGKA
l						AKAQREIEHSAGGPALLGDSVH
l						PPQPLARVLSPPLPGASRACWL
					ł	LRVRGPPSPRPPGTPAGPQAPH
		1				AAPVPARASPSTPPLVHLTNDV
	1				1	LRVKVKWAFFFTLHK
24056	54424	Α	24194	192	818	SWCLSRAVVGDSMTSEVIED\E
						KQF\YSKA\KTYWKQIPPTVDR
l						HAWGEGPNKTGTSCALDCGAG
ŀ						IGRITKRLLLPLFREVDMVDITE
						DFLVQAKTYLGEEGKRVRNYF
ŀ		1				CCGLQDFTP\EPDSYDVIW\IOW
ĺ						VIGHLT\DQHLA/ESFLRPCKGQ
						ASAPNGIIVIK\DNIA\QEGVILD
	1					D\VDSSVCRDLDVVRRII\CSAG
	1	1				LSLLAEERQENLPD
24057	54425	A	24195	h	233	FFFFETESRCVARLECSGAISAH
24037	34423	'`	24173	1	233	CNLCLPGASNSPASASO/CSWD
			1			YRCVPSRLANFLYFSTDGISPC
						WPRMVSNLLTS
24058	54426	A	24196	240	476	LLPQYRRVLLIYSGIQLLPGLGL
24036	34420	^	24190	240	470	RRRMCPGIYSFLLDFLVYLRRG
1						VYSIL*W*FVFVVLVPWFSAPS
						GHLRSSLCCLF
24059	54427	A	24197	288	677	NLPEDVVPDDKPHQPLLRAV\C
24039	34427	^	24197	200	077	LFVISKDLFLESPFESPIWRMYP
1						NSTVHIQRLELALLC/CHCPRAL
ł						PASAHSCHSLERGLPICKE**CM
		İ				PCSEYTIASQRPAGGAPAILPEC
1						DSRRRTTOATSEPGSHI
0.1000	54420	١.	04100	00	200	
24060	54428	Α	24198	99	399	LLCHGFEHPPLAQRSLLLPIV*S
						LLLSTRQSHSPSSFVPLLARSCG
						SWVSPAEAAEPQILQKGKCSCL
		1				IVPLEALCQRGTRLSLWLPLTFF
		_				PSFQLWPV
24061	54429	Α	24199	1	586	LSFFKVFSFFAIGSNILFSSEKFII
l		1	1		l	TDHLKPSSPTCQSHSLSSFVPLL
						ARVAFRWGFGVDVLSFVSFPSN
		1			1	SQDPQLQVCGVCWRSTPDRLP
l		1			l	GYHQRGSGGCRTADIIEPQMLR
					[LIVRLEVLSQRSTRPQRCQ\SA\L
					1	LGGASQLGCC/GVRDLLEEAVC
		1			1	PFSDL/KLHAGKTTTLFKAVRQ/
1		1			1	SHLSLQRLLLSFVCLCPT
		_				· - ·

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24062	54430	A	24200	11	822	
24063	54431	A	24201	1428	1801	RSAGIDPE*FFSG/PRRIHTASCL
24003	54451	ľ	24201	1420	1.001	PSQRSSNRACSSSVTRIVSILSRF
						IGIITPMNRNPRFWRNSTSWTR
					i	MNRQTSVNRFTTTLMSFTAAA
		1			i	SRVSVMTVKTSDTCSSRRRSQL
						VKYHLSISTQVL
24064	54432	В	24202	1	2040	THE THE STATE OF T
24065	54433	A	24203	176	564	AWEPKWYSTTATRTHFWEMRS
		-		1		RSIMSTTKAFSTSNSASDSLEEP
		ı				SSRKTRSSSLLQ*KSCNGGSEM
						SSTYRVVGLTGASLSASALNSLI
						DSRSNR*V/FSRNTLANITRRNO
						LKMLPAPDPDGFSSGSPFQ
24066	54434	Α	24204	I	774	
24067	54435	Α	24205	1	1807	MPLRAFNPVSSFRWARGMTIV
		1				AALMTVFFIMQLVGQVPAALW
		1		1		VIFGEDRFRWSATMIGLSLAVF
		ĺ		1		GILHALAQAFVTGPATKRFGEK
		l			i	QAIIAGMAADALGYVLLAFAT
		1		1	ŀ	RGWMAFPIMILLASGGIGMPAL
		1				QAMLSRQPNFDHWTADRHGD
		l				LCRLGEHMERVGMDCRRRPIPC
ļ		l				LPPRVASRCMEPGHLDLNGSRR
		l				HLANGFTTPRIGANQFLRRTVN
1		l				AQTNPWQNISIASAISSSRTRRIS
		l				GSVGSWPRSGNAEVSALHHYV
		l		ļ		PDLHRRMLLATLWNTYICINEA
		l				LALTLSDFSLVPPHPYRQLFTLT
		l		ł		TFHNRACSSSVTRIVSILSRFIGII
		l		1		TPMNRNPPYTEASVTKQEKTAL
		1				NMARFIRSQTLTLLEKLNELDA
		1				DEQADICESLHDHADELYRSCL
		l				ARFGDDGNSYDHDYAKLACLQ VEGRGGGTFAHELLAVEYAGW
		l				ISPAFRLKVMHWLMNRMLDAR
		l				NRDRYKQWKQDFFHDSYNRQS
		l				AGHILSQCANLAATTSEYFIHKF
		l				HRLIAAETGYSQSTVVRAFREA
		l				VNKGILSVEIVIGDHRERRADL
		ı	1			YRFTPSFLVFAQQAKNALIESKS
		I	I			KI\SSAATKVRLLAPISPTSPMG
						KIGLATSGS
24068	54436	Α	24206	291	851	
24069	54437	Α	24207	2	694	
24070	54438	Α	24208	2	331	HLSTIEPEDYNLLFRRVSLKLSV
			1			WLPTWQSKSNGWSKKCLWMH
1		1				LKSNPLNALLPPAGF*SATMPK
		1	l			CSAWAKTISSPGCVITEF*SQPV
		_				NAGTSPNKNTYLVGISPLKKP
24071	54439	Α	24209	1	4455	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24072	54440	A	24210	698	1088	LTSKFIAAVCMRRISLCGKGSA
						QNTDSGIAATPCSCVSD*QNSTS
ŀ	1	l				SRIRM*PWMCPCPMKRKSAT*N
		l		i		ITLRC*RAAPPISSRVRK*/MSM
l		l				Y/NHLQSAECGGHRPFPVPQEE
						SILLLHLIANRSSACHSIR
24073	54441	Α	24211	1	1902	
24074	54442	A	24212	1	2898	
24075	54443	A	24213	1	2604	
24076	54444	В	24214	1	1254	
24077	54445	A	24215	1	2946	
24078	54446	A	24216	698	1285	RFWRNSTSWTRMNRQTSVNRF
						TTTLMSFTAAASRVSVMTETA
		1				MTMITPSLHACRSKAVAAALLP
	ŀ	1				MNCSLWSTQAG/CSP/DVSAEG
		1				KPDIYRLSNRKITTCYSAES/LL
						KLSVWLPTWQSKSNGWSKKCL
						WMHLKSNSLNALLPPAGF*SAT
ł						MPKCSAWAKTISSPGCVITEF*S
		l				QPVNAGTSPNKNTYLVGISPLK
		l				KP
24079	54447	Α	24217	1	1986	
24080	54448	A	24218	1	2262	
24081	54449	В	24219	1	1014	
24082	54450	Α	24220	1	1155	
24083	54451	A	24221	5	386	
24084	54452	Α	24222	1922	2146	SSSAGISCWSSSPTPGSIPSRVGS
						AAA*GWTTSRSSTGSANPSASR
						KPAAAIRASMPPNCRSGEARWP
į						LWSGSL
24085	54453	Α	24223	1091	1435	
24086	54454	A	24224	1	2919	
24087	54455	Α	24225	1324	1502	
24088	54456	Α	24226	1	2901	
24089	54457	Α	24227	5	253	
24090	54458	A	24228	1227	3031	

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
24001	E4450	A	24220	1	1940	MNROTSVNRFTTTLMSFTAAA
24091	54459	A	24229	1	1940	
		1				SRVSVMTVKTSDTCSSRRRSQL
						VCKRMPGADKPVEGRGGGTFA
		1				HELLAVEYAGWISPAFRLKVN
		1				QTFIDYRTGRLQPAIPQSLPEAL
1		1				RLAADLAEQKQRLEQKMLMD
		1				APKVEFAERVATASGVLIGNYA
		1		1		KVLGLGQNYLFTWLRDNGILIA
						TGERRNVPKQEYISRGYFTLKE
		i				TVIDTSNGSRISFTTRITGKDVV
		1				RTTSIVMLAKVTFLSCITMSDFT
		l				FSGYELACFVTHSGLSRSAGHIL
		l				SQCANLAATTSEYFIHKPHRLIA
1						AETGYSQSTVVRAFREAVNKGI
1		l				LSVEIVIGDHRERRANLYRFTPS
		ŀ				F\LAFAQQAKNALIESKLKISSA
1		l			1	ATKVKAVLAKTLALFNFLSTPP
1						CQNDTPSPCQDDVAIKNKKSQ
		1			1	VKKTKRSVSGGAGTTSLKKLTS
						WIAKAKAKADNLRLSKKRTQK
						HEFKQKVEAAARKYAYLKNKR
		l				SPDIGGISNFDNLPHCMTVNEA
1		l				LNAVLAKNKDNEQWGAVAGA
		ŀ				YIADITDGEDRARHFGLMSACF
						GVGMVAGPVAGGLLGAISLHA
1						PFLAAAVLNGLNLLLGCFLMQ
1						ESHKGERRGELLKIVVLPGDHV
ļ		l			İ	GQEITAEAIKVLKAISDVRSNV
		ŀ				KFDFENHLIGGAAIDLQGGGGK
24092	54460	A	24230	1	1230	PRCVSKSLMLHCTR
24093	54461	A	24231	1	799	MKPRTYTVSVTAVKDGVSGVC
		1				SFRCSDVSRVSSFOWVRGLADF
		l				RSEAADLHNSGAQLASPIGSRT
		l				GATSGAPCQSRTHAPALLSLW
						AVDGTGRHGAGGSTRRQGSGG
		l				TQEPMVAGKAQAWRAAGFSG
		ŀ				SGDGVGVSILVPWGGTCIGSFR
						OANSWISGGAFSDAGCDSDRL
		ŀ				GKDWNSLEGSEEERKMWESLE
ļ.						LPRDLKSSEDRKMWESLELPRD
						LLNGFDQNAD\TICTMKSRLRW
		l				SQMEMKNLLGTGVNVTLVML*
						QRDWRHLLLV
24094	54462	Α	24232	416	775	
24095	54463	Α	24233	548	882	PGGHQYPPCHLEEKETDLPLLY
						TAVEEKGLSYSIIPGSLVEKPLT
				l		STNSSGTPIQSNITSL*IPLAGIMS
		1		l		MAASQN*AI*WPLAPTMQSFTC
						TT*RLS*SHRRSWWHRGCT
24096	54464	Α	24234	144	570	

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	sequence	11100	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			,	sequence		, , ,
24097	54465	A	24235	254	1117	LSRMKLRTLVVSVTVLKDVVS
						GVCSFRYSDVSGVSSFQWVRG
						LADFRSEAADLRGLKLOTFVVS
						VTPHKGSVDPKSDQQRDLLQR
						VKEONFRNVKGDPAALAHEDD
				1		VNGTESGAPRKHGDRELDDVV
			ł			ITNPHIEAILENKDWIEDASGLM
						SHCIAILKICHTLTEKLVAMTM
						GSGAKMKTSAIVSDMIVVAKW
						NSPRVDDVVKSMYPPLDPKLL
						DTGTTTLLLSV\SHLVLVTRNA
						CHL\TG\GLDWI\DQSLS\AAEEH
						LEVLREAAL\ASEPDKGLPGPEG
						FLQEQSAI
24098	54466	A	24236	472	844	RTDGVLVWMSFLFVSFPSNSQD
						PQLQVCWSLLEVHSRPCLPGYQ
						QWRLLCPHEITGESMSKLHIIQK
		1			1	YTSSCCLFKSS\SERGSSTPCPSW
						LGSGEGLSTLHATDFSFSAPMA
						QRGKSSLVSLE
24099	54467	A	24237	206	456	
24100	54468	Α	24238	50	392	LGLLPFFHRCPTQRGGI*RGSLA
						TAALLSCGGLCPVGS\SRHLCL\
						PVRGKRPTQASVMGDALPPTK
						LECPRSASGCCAGSDNFKPVDL
		l				SLLSSMGLGSTELDHLAPWLQP
24101	54469	A	24239	1	204	PFQG SFSVOFCSLVGDPLG*SSCDPLE
24101	34409	IA.	24239	l'	204	EKRCSGFWNIQPFCAGFSPSLW
						DYLSLVFDVDDLQMRSLSGCPF
24102	54470	A	24240	1	375	MCNAQVLEYMGKSSSLLTSDL
24102	34470	ľ	24240	ľ	373	OLVRDALRSLRNSFSGHDPOHH
		1				TIDSLEQGISSLMERLHVMETQ
						KKQERKPPLVIPRQT/WVWSGP
						PANSNRPAAEGPDC*KEN*RDA
						LRSLRNSFSGHDPQHHTIDSLEQ
						GISSLMERLHVMETOKKOERKP
				l		PLVIPRQTGSGVDLQQTPTDLQ
				ĺ		LRDLTVRRKTNEOKGIASASTK
						RTSTPKPHL
24103	54471	Α	24241	1	674	MVVYACSPSYLEAEVGGSLSLE
						VEATNPKNSSKKFLDLISEFCK
						VSGYKISVHKSVALLYTKNDQ
						AATQIKNLIPFTTAAKQFKYFGI
						YLTKEEERLKPSRKKITKKHTK
						KRTASLILHAMICSQIPKQQQNE
			i			KYQVPQFDQSTIKNIESAKGLD
						VWDSWPLQNADGTPPLVIPRQ
				ŀ		T/WVWSGPPANSNRPAAEGPDC
						T/WVWSGPPANSNRPAAEGPDC QKEN*QTERNSINIKKKDIHTKT

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
24104	54472	Α	24242	1676	1768	
24105	54473	Α	24243	270	471	PSREYQDANPWLKWIKYSVRT
		l				LNKSDCYASGYK*ACLLCSVYS
	1					HGKAAGECTLSAGSKNGLAEVI
24106	54474	-	24244	23	200	KK AOMKAHGRSRSS*RC\DEEHWR
24100	34474	l ^A	24244	23	200	ACRAQEPPCSQSQRERSKRTCR
1		1	ļ.			QGTVASKVKCHSEAR
24107	54475	A	24245	3	270	ASTSHVPQPICGLDDLAIRHSAV
24107	34473	ľ	2-12-13	ľ	2.0	RLNRERVHSTAVLP*EYTKQRR
		1				QGHL*PDPYKKEGAATNEDAL
		1				LLRWERGRGARLHIPGCKAPQL
24108	54476	A	24246	135	275	
24109	54477	Α	24247	365	583	
24110	54478	A	24248	2906	3114	
24111	54479	Α	24249	108	405	NLHTPYRPQSLGQGEPLDDYV
					ŀ	NTQGAS/LFSVTKKELGAGSIEE
1			1		l	CAAKCEEDKEFTCRYFHCRCTF
		1	1	İ		LSIPQVTSSFLCFPLPLPLLKSSK
L		Ļ.				LLANRDKYRM
24112	54480	A	24250	1	386 808	MSYSOCVSDPKAHGILPGYHC
24113	54481	I ^A	24251	1	000	WLFRDQGLLKSQHVINPAKAST
		ľ				GFFLODTLMVLEIKTTIRNYYK
		İ				HLYAHELERLEEMDKFLDTYT
						LRRLSOEVIDSPNRPIMSSKIESV
1		l				KGYTHOOFCHKSTLNKGDRVI
						DNLTCPPYCCVWFPLAGTGPHI
						LYLSRLVSNLELFKRGKGRGEQ
ł		1				RKEEVTCGMLRKSLVTSDLPEP
		1				GGVTSEGCETAKVTACSSLWE
		1			1	LCHREVLTPYCPKDTCRKWLE
	1	1				TPVRRSCPVRRSRISTCLRKKFG
						HILAEQLSCA*RCL**FL/YGCFY
		ı				FQDHECVLKEESSTSFGRINHV
						LTLKEPLVPE
24114	54482	Α	24252	833	1245	ANMFPAFAAVAAGAATEAAA
				1	1	AAAAAPAPLAPARRAAAAARG
		l		ł		PPLPRLP/WSCAEVMYHPQPYG
						ASQYLPNPMAATTCPTAYY/RA GAPTWPAGESPDGQPAGGGGE
						GHREYRSGARRRPAGTRGAED
				1		ALGSAGASLSRKP
		_	L	L	L	ALGOAGASLOKKY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	i	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
24115	54483	A	24253	145	766	TGCIREKFHLPEGSLLIVSSRKH
		ľ.				LORLAFOAKVSSMMLPLOGAO
				1		MLOMLEKSLR\KSLPRNPLKVY
1						GT\VFHINHG\NPFNLKAVV\DK
						W\PDFN\TVVV\CPQ\EQDMDRW
						TLDHYTIT\YOIYSKDPONCOEF
Į.		1				LGSPELINWKOHLOIOSSOPSLN
		l				EAIQNLAAIKSFKVKQ\TQRILY
						MGRLETAK\ELTPFLAGNQRILS
						P\SGWAKPKAM
24116	54484	A	24254	215	550	VGFRSCLVMSGRGKGGKGLGK
24110	34404	^	27237	213	330	GGAKRHRKVLRDNIOGITKPAI
					1	RRLARRGGVKRISGLIYEETRG
				i		VLKVFLENVIRDAVTYTEHAK
						RKTVTAMDVVYAL\KROGRTL
24117	54485	A	24255	3	292	TLLNLAAAHCSGPCLLWFOLSF
24117	34403	1	24233	ľ	272	RSPSTSAVGRCRRPSADFHPSRS
						SRVSAVLLIQ/QRCPLPILIQLKA
		l		1		CHCSCTAKWLGSSPKPETFFEII
						HPSPS
24118	54486	Α	24256	1	789	111 51 5
24119	54487	A	24257	3	256	TLLNLATALSSGPCLLRLELSFH
24119	34407	l^	24237	ľ	230	SPSTTAVCCRHRPAADFPPFGR
		l				VSSLLLIQ/QRCPLPLLIGLKACQ
1						CSCAAKCCVRPNRAEQ
24120	54488	A	24258	3	212	CSCHARGE VIG MOLEQ
2412I	54489	A	24259	2	491	
24122	54490	A	24260	20	272	
24123	54491	Α	24261	160	876	
24124	54492	Α	24262	623	796	GANSWNLTGTLAVKAGTVENP
		İ				I*VFATHVESAGPR*LDPRTGSL
		1				AVRTKGRHWDQGA
24125	54493	Α	24263	242	659	TSNSPPOTRPHVCHGPRQYLLQ
						LSSRDLEFLVQIAVRSCLCPGLR
						IRSPVFAVASTSTFGTLRGKVV
						KDFGFVFPPPWGQPCLCPVEER
İ	ł					ALSCOGPVVDSRCPRCGHEGM
						AYHTQQMRSADEGQTVFYT\C
		1				TNCKFQEK
24126	54494	Α	24264	66	238	,
24127	54495	Α	24265	2	278	
24128	54496	Α	24266	168	565	TENFRALSTGEKGFGYKSSCFH
					1	RIIPGFMCQGGDFIRHNGTGGK
				1	1	SIYGEKFDDENFILKHTGPGML
					1	SMANAGPNTNGSQFLICTAKTE
			1		1	WLDGKLVVFGKVNGMNIVEA
						MECFVSRNGKTGQKITIA\DCG
24129	54497	В	24267	20	88	
24130	54498	C	24268	183	254	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, v=possible nucleotide insertion)
24131	54499	A	24269	412	590	KSGLATQG/GRRRGWVFLAPQ KGGEGVETQREGVGVLAPSPE
						KRDLPLRVKDQGRRPCVV
24132	54500	A	24270	550	852	
24133	54501	A	24271	207	381	
24134	54502	A	24272	412	589	KSGLATQG/GRRRGWVFLAPQ KGGEGVETQREGVGVLAPSPE KRDLPLRVKDQGRRPCVV
24135	54503	A	24273	191	235	
24136	54504	Α	24274	95	375	AEKIWEGVSQRALGQSSRGSGS GCQMNPLSGRPQVVCFFTAHM LTKKREEGASMLNMPCFQ*KE EMIQANLGPIADSSHCLGIEGPP WAPLTP
24137	54505	A	24275	144	293	WQVPLFWQPFPEPLELWPKAL* LTPSQIFSD*RLKTDAARSPRKP PAPR
24138	54506	С	24276	1	909	
24139	54507	A	24277		1744	MGCLGTORAPRAFLLPLPLHE SGGGGERGATAAGTEQGAC LAASLLRGLWDAGEVTFIRVR SRPVDPTLAPLHEHSEAQCVAS PSLTVTKFKSLLPKSLLAGSRE PSLTVTKFKSLLPKSLLAGSRE RSRALLFOQARPPREPROVTQ RTQLQGQRILQWRPATEGPHSP RRGSDSAKGTHPPTTGTCPPRR RQGAGYSRYNSGESRRLGRQ QDLVARSDEQRDPAPATRRED KMIEELQPFGEHRPRRSPSGQDL VLALCKAAAGPGAPPAFRRED GDTLFQALWFLASPSFQRGQLC VLALCKAAAGPGAPQASTAG ADLRAFMGLRGEVCADWSM GGHGSGECRGTGGQRAVVK SPVQEKGKEPLSLLQYFMVDEB ERFLRITISHEDNDNSNSSNHS SINETKETCFHEPKTPAPVTDW EGSLLYFNNCRDASLHISCFSK SWCRGKYPSTSPSPSPLASPTFL SWGRGKYPSTSPSPSPLASPTFL SWGRGKYPSTSPSPSPLASPTFL SWGRGKYPSTSPSPSPLASPTFL GQGGELVTSRNATUR GPGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ GGGFLLSHVPSVRDPTGNTVQ
24140	54508	A	24278	2	120	LTWQPLPEPLELWPKAL QSSAAKPRTSGKGSVREPWAR
-7170	15 7500	1	1-72.0	Γ	I	VPGALGVAAR*AEQSDFQ

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /-possible nucleotide deletion, \-possible nucleotide insertion)
24141	54509	Α	24279	1135	1483	ILTMTGSLSCTPSNAVIEKV*VC CSSLSTG*SKVQVASNTSLCDG ED*GVVIQKKVKLWVFCIFYVL VQSYYS*R*SQIETVRNGCSHW GRFQDIALITGHCKIWTLIVFIQ YFNV
24142	54510	A	24280	1	860	ATTGGTNLRDDMRLDDTVHU VIATPGRLDLIKKGVAKVDHV QMIVLDEADKLLSQDFVQIMED III.TLPKNRQILLYSATPLSVQK FMNSHLQPKYBINLMEELTLKG VTOYYAYVTERQKVHCLNTLF SRLQINQSIFCNSSQRVELLAK KISQLGYSGPYIHAKMRQEHRN RVFHDFRNGLCRNLVCTDLFTR GIDIQAYNVINFDFPKLAETVL HRIGRSGRFGHLGLAINLITYDD RFNLKSIEAQL*TEIKPIPSNIDK SLYVAEYHSEPVEDEKP
24143	54511	Α	24281	848	1218	PMQRKSMLKISVS/PVSVQKFM NSHLQKPYEINLMEELTLKGVT QYYACITELQKLAETYLHRIGR SGRFGHPGLAINLITYDDHFNL KSIEEQLGTEIKPIPSNIDKSLCG AEYHSEPVEDEKP
24144	54512	A	24282	241	477	
24145	54513	A	24283	38	208	MTAGAPPPPPPFET\PEGV*GNP TCSGG*EDASGSTRSQRHPVGL PVSCALRACHLK
24146	54514	A	24284	794	1455	AQQGGTSKKRRFPODGOQRAG LACSRPPOPGCGCFQAPAGERY PNIKPHPTSVAPLVLCICYLMP MLPWAWSYWRGARRPSRHPG AFPCSVAHQCPSAARSPTLRQE DPEKGSQASSFPTGSPOGMVD RCLKGPQLDVSRKQLSLPQTGL RPVLVNQLSKKPTGYPQTQIQW VLMRDTGKDTDTEEKEGHAK MEAEIGVMYPQEHLEPPESGRA KGEIPL
24147	54515	Α	24285	90	354	QSPGVHQLSRFVS*SYSERP*SL LVSTRYHTRALH**HYADSIQ* ARSSKHTGLIGCTVP*RGFPRTH SFYSRKREPRRQTSSFRGIL
24148	54516	Α	24286	236	520	TACFLTHVAIKWVMHSSIPSSN GGGIYMIGLKAQVSYMRKWLK SPWSPLLLSPSLHRWPHGKFPM IS*QRKRRLGPGSQMVLHDMQ APPKSGQL

SEO ID	ISEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
110.	sequence	liou	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
1	acquence		057540,217	sequence	or prepare sequence	
	ŀ					
24149	54517	Α	24287	16	278	FAQAMKPHLVQQLQLESPLG*
		1		1		AYDNPLSFSKIHLSDLHKPVL*L
						GDSNDVLGTLESMSAQSQCPV
		1		1		VPEMSDHLPFPMHSYPGKRPEV
24150	54518	A	24288	74	168	LGKCLFLHSCP*APPEAICLQLP
						LLSSPNGP
24151	54519	С	24289	92	564	
24152	54520	Ä	24290	1	876	
24153	54521	A	24291	71	282	PLVTQVVFRLFPTNSSLDLLQ/S
						PSP/GSGPHSNWQPFGSLDKWA
1	ļ		1		Į.	GVKHPNEECAASKIQTGPPCVV
		1				PLPWL
24154	54522	A	24292	81	1054	PRLFSDCFPPTAAWTCCRAFSC
	i .	1				SGPHSNWQPFGSLDKWAGVKH
					1	PNEECAASKIQTGPPCVVPLPW
						L*EEPNAAIYPSPVSPVCFLLLA
	1					HWIOSA*CHOCNGPVWYLVEV
	1	1				KSNQVLSK*HYDTKPES*YTPY
	1	1				GOEWRKGHLPSOWPHTRYOE
1						MY*FAQAMKPHLV*QLQLQSP
		1				LG*AYNNPLSFSKIHLPSPHGSH
1		1	l			SEPCL*GPTRTLV*L*L*KQTGM
		1				LAVAPEENQHYLAWQLPQGRP
						SLLPOAVOGLSPOTKVERLMA
						ASVGEGMLPILGMEKLF/PSGK
1	l					KD*SGFTSSVSPASSGSSHTSPF
		1	1			QVAGSHSFPINAFTLTVDTW
24155	54523	Α	24293	5	88	`
24156	54524	Α	24294	779	1041	PPLSSPSGP/DEQSD/LWWQQSIP
		1	i .			LHQETVKVLH*MGPVPCATQL
						DKIHOOGFSDPYRGAVLLASG
1		1				WCPKRPEIPEEGATTHLCCFPAS
24157	54525	A	24295	248	3714	
24158	54526	A	24296	4177	7804	TFQESRFEQGFQKAIFILELVDA
						EDPPDEEEEASSIFSSSFHFLFP
	1	1				SSSSCFSFSSSSSSLLLRSPGDKD
		1				MPTAGMPSLLQSSSESPQSYPE
		1				GEDSQSPLQIPQSSPESDDTLYP
		İ			1	LQSPQSRSEGEDSSDPLQRPPEG
1			1			KDSQSPLQIPQSSPEGDDTQSPL
1		1	1			QNSQSSPEGKDSLSPLEISQSPPE
1					1	GEDVQSPLQNPASSFFSSALLSI
			1			FQSSPESIQSPFEGFPQSVLQIPV
	I		1		1	SAASSSTLVSIFQSSPESTQSP
24159	54527	A	24297	145	219	5. 2. 2. 2. 3. 1 5. 5. 1 Q 55. 2. 2. 1 Q 51
24139	34321	1^	27271	1175	217	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	delction, \=possible nucleotide insertion)
				sequence		
24160	54528	A	24298	191	1075	OPKVSRHCOTSPGSYLYSLOSG
24100	34328	^	24230	1127	1075	RWTTGKPLQQTKSVYLRSFHH
	ì					HHFONHRHYLHHHHLYHHHH
						YHPHPSHHHLHCY\SOHHHHH
	İ					HHHHHHHEHHLCRHOKSTOVE
i			l			YSVFAERENDLGLNGTGDEAL
						HDNGTTERVSGPPAGYLYRAD
						YAIGLQPFMPRLGRRTPGAGGP
						HSRAPELAYLQKLITSPATLPFV
						YLETGGWSPRDAHRVDNFDM
1						GPRENMGSVWEKSFLFTRNPSK
	1				i	SSYIADKNLIYLYASECIGRQSQ
	1				i	KARASKSGVQAEHPYITGAIKD
		L				DTRAIVKTRDLANRLK
24161	54529	A	24299	155	381	LICTSFKILSRLFCPSSL*SSSSQG
	1	l				ACISKLLSVLTYAHLYFALEKE
-	ŀ					VLTFLSRAIVSSSTLLTHGVAHS
						LTLFH
24162	54530	A	24300	1052	1338	TGWDGPPANCSSPMEGDSNVH
	1			l		CSCVCCCSCFGLLL/CAIHCCCA
	1	l				PPGCSACGAATLLYSAPLGSSA
	1	i				YGIALLSPFHHVRMMLSTKQEV
		i				GLHQTLNLP
24163	54531	A	24301	I	723	MLNVQALVVMVMCFMWASL
						WAPGGHPSAGNISSWVSGLVSF
					ĺ	PEWSSWELVVSVVGWAGKLW
			ł			LHESLLTLNLSCGLLVVLVALV
						FDEWHPWSGHVAEACSTGYWS
		l				TGGIWFHEIAAP*DCSSSPAME
	İ	1				QSWMEND\DELIEVGFRRLVIT
		1		l		NFSELKEDVRTHLKEAKNLEKR
	1	1		l		LDEWLTRINSIEKTLNDLMELK
	1	1		l		TMARELRDACTSFSSQFDQVEE
1	1	1	l	I		RVSVIEDOMNEMKREEKFREK
1	1	1		l		RVKRNEQSLQEI
24164	54532	c	24302	295	430	
27104	107002	<u> </u>	24202	270	730	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	coden for last amine acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
,				sequence		
24165	54533	A	24303	811	1947	OTERNSININRNGHOHONPICRS
2.1103	15.555	ľ.	2.000	l*		PSSKTKGSIKPORSGRNOSRKA
				1		ENSKNOSTSSPPKDCSSSLAMK
			1	1		OSWMENDFDELTGAGFRRLVI
l				1		TDFSELKEDVOTHIKEAKNLE
1				1		KKIRQM\VTRINSVEKSLKDLM
	i			1		ELKTMARELCDACTSFSSRFDQ
		1		1		VEERVSVIEDOMNEMKRKEKF
		1		1		REKRVKRNEOSLOVIWDYVKR
ŀ				1		P\NLRLIGIPESDEENGTKLENTL
						ODIIQENFPNIARQAN/VOIQEIQ
				1		RTPQRYSSRRATPRHIIVRFTKV
		1		i		EMKEK\MLRAARQKGRVTHKG
			1	l		KPIRLTADLSAETVQARREWGP
	1					IFNILKGKNFQPRISYPAKLSFIS
		1				EGEIKYFTDKOMLKDFVTTRPA
				1		LQELLKEALNMERNNWYQPLQ
24166	54534	Α	24304	 	2394	EQUEENTER WILLIAM TO FE
24167	54535	A	24305	1902	2413	
24168	54536	A	24306	2	610	WAPPAIHIRCMPEAYNFPRKPL
		1				SLIPSAIQSNELIIMNGNRALNIQ
						LLKRYOCTLDIRYKOTSMLNC
				i		YEGTVITPSHIOYPCLSDSYGLY
						DIGIRNTDFLKGELSMNIRPLHD
						RVIVKRKEVETKSAGGIVLTGS
ŀ						AAAKSTRGEVLAVGNGRILEN
ŀ					,	GEVKPLDVKVGDIVIFNDGY\G
				1		VKSEKIDYEEVLIMSESDILAIV
24169	54537	A	24307	I	801	
24170	54538	Α	24308	1268	1579	TAIWSPSKSALYAAQTSGCSWI
				i		ALPSISTGSNAWIPRRCRVGARF
			1	1		SSTGCSRITSSRISQTTASSRSTIF
	i	İ		1		FAALMVVARPRSSSLP*MNGLN
1				1		SSSAIFFGRPH
24171	54539	Α	24309	583	796	IHICDIRFSATPAGGEFWQYIAS
		1			1	RQSIFHAIVETGHALIQPSSVPG
		1				LTPAMKAVATTSH*TVESCFRS
		1		ŀ		ASR
24172	54540	Α	24310	I	313	MPNVDVVGEMVNTMSASRSY
		ı		ľ		QANVEVLNTEKAMSIAVTTTDP
l						TNTGVSTTSSSSLTG\SNAADLQ
l		1				SSFLTLLVAQLKNQDPTNPMEN
			1	ļ.		NELTSQLAQISTTLRAY
24173	54541	Α	24311	54	323	SASAQVSALSIHSAAFQNEYLSP
1		1				CPN*RLV\SAFTNSSRQSG*PEK
1	1	1		l	1	SVSQTPVITASAFT*YADIAASV
						RNKILRPGTNVFGDTIGFRVQ
24174	54542	A	24312	816	866	`-
24175	54543	В	24313	1	985	
24176	54544	A	24314	1828	2505	
24177	54545	Α	24315	1	1938	

PCT/US01/08631

WO 01/075067

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
24178	54546	Α	24316	420I	4668	
24179	54547	Α	24317	269	503	
24180	54548	В	24318	l	1702	
24181	54549	Α	24319	829	1584	
24182	54550	Α	24320	1238	1926	IQAAYNRIAPSSSPKFPRACGKP
	Ì			!		CLGERGRIRGPGPSFGLINRPST
				l		PDPTITGVSTT/SSSSLTGSNAAD
	ł	l		l		VQS/SFLTLLVAQLKNQDPTNP/
	i			İ		MENNELTSQLAQISTVS/GIEKL
		l				NTTLGSISGQI/DNSQSLQASNLI
		l			1	GHGVM/IPGTTVLAGTGSEEGA
		l		1		V/TTTTPFGVE\WDGTL/TDGST
		l		i		APNGSYNVAIS/ASNGGTQLVA
		l		1	ł	QPLQFALAAERFCALSPGAPYR
		L_				YDTWVRQWVASP
24183	5455 I	Α	2432I	487	1509	FYLQGLRQQPQVRQPLNIGVTA
		l				QGVNAAACHPHIAEQQLDHRH
		l		1		RTNVLRTNGVLRPAEGIQERGS
		1				FIVSTGFCDVFADFQERVFWRT
		١				TDVFNHIWRVAGNVLFQQVPH
		l				AARMLQRGIAFGEAVFVQFISP
						GGFIVLAFFRVVAGEQAIFKAVI
						LTHDQAGGGIGFGVFAVEFFIV
						QQVQQYA*QESNVSTGTNRGIV
		1				VSYRRRTRKARIDHNQLRAVV
		l				NFGFHRPAEPDRVSAHHHDEV
						GVLDVDPVVGHRKTFFDIWHD
		l				DQFIDQRVRRFSGDNRRLQDRE
						QVGLQSANNSPIDKSLFRGINQF
		1				QLYAAFTAQDVNIDIFKTRQQF
						FAVIGQTTGVQYGK
24184	54552	Α	24322	377	749	
24185	54553	A	24323	140	298	
24186	54554	Α	24324	342	1094	MGAPWLHRPRRALQRWRLGS
		l				GFSCGSGLGLQENNGGRPVMV
		l				LALSTLPRRPTRCGCEFYCLIKP
		l				GFRLGMKSVKMEIRGLWAGGE
	l					DKIAVPAEPRSAGPERTLSDLH
	1					NESIF/ITGGGSGLGLALVERF/IE
		l				EGAQVATLELSAAKV/ASLR/Q
		1		1		RFGEHILAVEGN/VTCYADYQR
	1			ŀ	1	AVDQILT/RSGK/LVNTPAETLE
	1					TGFHELFNVNVLGYLLGAKAC
		1		I		APALIASEGSMIFTLSNAAWYP\
21105		ļ.	0.4225	l	1005	GGGGPLYTASKHAATGR
24187	54555	Α	24325	I	1785 1760	<u> </u>
24188	54556	A	24326	2		TEOMICOBIICDEEDHVARWAY
24189	54557	Α	24327	2437	2679	TFQYIGQPHGDEFPHVAPVKHG
		1				DKLGEGKNHHDKSRHLFHRFN
		1				H*MR*IRLTSNCAGSDAISQRD
	1				l	NKEKKNDQWHDTFNKG

SEO ID	SEO ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
ł				sequence		
24190	54558	A	24328	3	415	
24191	54559	A	24329	ī	1932	
24191	54560	A	24329	1932	2117	RAISLGRTAAGYLNSDVAQRA
24192	34360	^	24330	1932	2117	ASGADPAGDFVVWH**IRKNLS
1				l		
		ļ			1017	GGARYAIPHLYQHLAWHP
24193	5456I	Α	2433 I	1	1017	MRYREPPARGVEPNELHACRSE
						DKLSEDAFDDQCTGANPRYPLI
				l		SELKQILLDTYYGRDYVEDRLS
						FLLAVATLNGERLDGEMSEGK
			į.			LVMPFRHVSDAFEQTRETIGVR
				l		GNNAINDMVRQRLLN\RFTSEQ
						AEGNAIYRLTPLGIGITDYYIRQ
į				1		REFSTLRLSMQLSIVAGELKRA
	1	ļ	ľ			ADAAEEGGDEFHWHRNVYAPL
						KYSVAEIFDSIDLTQRLMDEQQ
		1				QQVKDDIAQLLNKDWRAAISS
	i	1		l		CELLLSETSGTLRELQDTLEAA
		1				GDKLQANLLRIQDATMTHDDL
	i	1				HFVDRLVFDLQSKLDRIISWGQ
1		1				QSIDLWIGYDRHVHKFIRTGID
				1		MDKKPRPSRRCL
24194	54562	Α	24332	5	220	GLSWLNYC*ASTLTISLRCATR
		1				AVPLTRIRCRPRLLPSRRERTAL
		1		1		PCIYVKIAVTLLTATCASCVRR
		1				WIPA
24195	54563	Α	24333	3	180	
24196	54564	A	24334	229	564	PGFKPRHSSWQRLPSLGHLPLY
						GSSVFTLLNLATAHSSGPCLLQ
						LELSFHSPSTTAVCRRHRPTADF
						HPSESGRVSHVLLIQ/QRCPLLL
						WIRLKACHCSCTAKCPGSSQLS
24197	54565	Α	24335	148	404	
24198	54566	Α	24336	213	1306	SCEHSVAHGRGIMDLSLLKALS
		1				EADAIASSEQEVRQIL\LEEADR
		1				LQKEVRVDGLGSVLIRLNESTG
						PKVMICAHMDEVGFMVRSISRE
		1		1		GAIDVLPVGNVRMAARQLQPV
						RITTREECKIPGLLDGDRQGND
						VSAMRVDIGARSYDEVMQAGI
		1				RPGDRVTFDTTFQVLPHQRVM
		1		1		GKAFDDRLGCYLLVTQLRELH
		1				DAELPAEVWLVASSSEEVGLR
	1	1	1	1		GGQTS\TRAVSPDVAIVLDTAC
		1		1	1	WAKNFDYGAANHROIGNGPM
		1	1			LVLSDKSLIAPPKLTAWVETVA
		1				AEIGVPLQADMFSNGGTDGGA
		1		1		VHLTGTGVPTVVMGPATRHGH
			1			CAASIADCRDILHMQQLLSAFI
		1	1		1	HKSNELGGVLRLRIQICRDGNE
24199	54567	В	24337	I	1075	THE SECOND PROPERTY OF THE PRO
24127	124301	10	~~331	1	1.0.0	

NO:	SEQ ID	ISEO ID NO:	Met	ISEO ID NO:	Nucleotide	Nucleatide location of last	Amino acid sequence (X=Unknown,
Sequence							
24200							
GDSVSTGLVVAHISHVANITIPIG		'			sequence		
GDSVSTGLVVAHISHVANITIPIG							
LYLLNPSSGADGKKNSNLSALI SAAKEPVVWAPVLATILVLVG	24200	54568	A	24338	438	1462	
SAAKEPVWAPQLATILVLVG			1				GDSVSTGLVVAIISIIVNAITIPIG
VKIPAAWPTFNIJAKANSGVA		1					LYLLNPSSGADGKKNSNLSALI
VFAAGLTLAAHKEEFSAEJAYN							SAAKEPVVWAPVLATILVLVG
TFI,KILIMPI,ALLLVGMACHLN SEHLQMINTLASALPPAFSLKLIY CAA/ARLMCYTRTGITSLATIT ETVTQKLTGSEPBISSYGRNLM WPLAWYKDNOKPHLCSQLETT RIHSTAAQGRSSGSLRRGLGAS PFYRQHDQARSEGQIPITRIPMG LAQRAPKGRAQPRGLTAFITM GALQQPQGRGTSKQASQTEPR ASAWPHTPT GALQQPQGRGTSKQASQTEPR ASAWPHTPT GALQQPQGRGTSKQASQTEPR ASAWPHTPT GALQQPQRGTSKQASQTEPR ASAWPHTPT GALQQPQGRGTSKQASQTEPR ASAWPHTPT GALQQPQGRGTSKQASQTEPR ASAWPHTPT GALQQPQGRGTSKQASQTEPR ASAWPHTPT GALQQPQGRGTSKQASQTEPR ASAWPHTPT GALQQPQGRGTSKQASQTEPR ASAWPHTPT GALQQPCGCHNNAASDRLRSYKK QPAER/WLAPPIDSITSPPASRRA LILLY L					l		VKIPAAWDPTFNLIAKANSGVA
SEHLQMITTLASALPPAFSLKLIY			1				VFAAGLTLAAHKFEFSAEIAYN
SEHLQMITTLASALPPAFSLKLIY		1	1		1		TFLKLILMPLALLLVGMACHLN
CAA/ARLMCYTRTGTTSLATILT				1			
ETVTQKLTGSRPRISSYGRNLM							
WPLAWYKDNGKPHLCSQLETT RIHSTAAQGRSSGSLRGLGAS PFYRQIDQARSEGQIPTRIPMG LAQRAPKGRAQPRGLTAPFTN GALGQPQGRGTSKQASQTRPR ASA WPHTPT GSGYQRCHNNAASDRLRSYRK QPAER/WLAPPIDSITSPPASRRA LILLY					į .		
RIHSTAAQGRSSGSLRRGLGAS	1	1					
PFYRQHIDQARSEGQIPITRIPMG LAQRAPKGRAQPRGLTAPFTN GALCQPQGRGTSKQASQTRPR ASAWPHIPT	1	l			l		
LAQRAPKGRAQPRGLTAPTN GALGQPQGRGTSKQASQTRPR	1						
GALOGPOGROTSKQASQTRPR			l		Į.		
ASA WPHTPT					ĺ		
24201 54569			1		1	1	
QPAER/WLAPPIDSITSPPASRRA			_				
24202 54570	24201	54569	A	24339	30	177	
24202 54570			1				
24203 54571							LILLY
GACLTSSSCRKGQLARRGLSYQ SPHAPADPAPAARRSPYCLS			Α				
SPHAPADPAPRARRSPYFCLS DDGGMKSPRDMSPDKPSRHPD QVALTRRIPSPPAVALPSVVKR TVHEGQGDDDESNPSPLSPDVH	24203	54571	Α	24341	425	824	
DDGGHKSPRDMSPDKPSRHPD			l				GACLTSSSCRKGQLARRGLSYQ
QVALTRRIPSPPAVALPSVVKR QVALTRRIPSPPAVALPSVVKR TVHEQQDDDESNPSPLSPDVH		l.	l				SPHAPADPAFPAARRSPVFCLS
TVHEGQGDDDESNPSPLSPDVH		ľ	l				DDGGHKSPRDMSPDKPSRHPD
24205 54572 B 24342 I 2118		1	1				QVALTRRIPSPPAVALPSVVKR
24205 54573 A 24343 I 888	1				1		TVHEGQGDDDESNPSPLSPDVH
24206 54574	24204	54572	В	24342	I	2118	
24207 54575 B 24345 I 1992	24205		Α				
24208 54576 A 24346 880 1103 TMMRRLISRÖPVLPLÜKCESFÜLI SNPWRVFGGTGGORÖER/CRY AEBSVAETRLEP*CRLYAEVDP** GSGCAGLSL** 24209 54577 A 24347 1 952 MITIKKLPLAVAVAAGVMSAQ AMAVDFHGVARSGIGWTGSGG EQÇCFQTTGAQSKYRLĞNECE TYAELKLĞQEVWKEĞDKSFYF** DTNVAYSVAQQNDWEATDPAF REANVQĞKNLEWLPGSTIWA GKRFYQRHDVHMIDFYYWDIS GPGAGLENIDVGFGKLSLAATR SSEAGGSSSFASHHYDYSSNEN ANDVFDWLARWDSFGGFLTG FPPSALACTSGVSG(GKPFGPP EAQWTFATPKKPGKEACSSIQV LNRGYRPPIFSYNTGVRRSARD YYPPRQRSISTNSPSRRWKGKL					730		
SNPWRVFGGTGGGNGENCRY AESVAETRLCP*CRLYAEVDP GSGCAGLSL 24209 54577 A 24347 1 952 MITLRILPLAVAVAAGVMSAQ AMAVDFHGYARSGIGWTGSGG EQQCFQTTGAGSKYRLGNECE TYAELKLGGEVWKEGDKSFVF DTNVAYSVAQQNDWEATDPAF REANVQGKNLIEWLPGSTIWA GKRFYQRHDVHMIDFYYWDIS GPGAGLENIDVGFGKLSLAATR SSEAGGSSSFASNHIYDYSNEN AND/YEDVRLARWDSPGGPLTG FPPSALACTSGVSG(KGYDGPP EAQVTFATPKKPGKEACSSIQV LNRGYRPPTFSYNTGVRRSARD YYPRRGRISITNSPSRRWGKLK					-		
AEBSVAETRLCP*CRLYAEYDP GSGCAGLSL 24209 54577 A 24347 1 952 MITLKRLPLAVAVAAGVMSAQ AMAVDFHGVARSGIGWTGSGG EQQCFQTTGAGSKYRLGNEGE TYAELKLGQEVWKEGDKSFYF DTNVAYSVAQQNDWEATDPAF REANVQGKNLEWLPGSTIWA GKRFYQRHDVHMIDFYYWDIS GPGAGLENIDVGFGKLSLAATR SSEAGGSSFASNHIYDYSNEN ANDVFDVRLARWDSPGGPLTG FPPSALACTSGVSGGKGVFGPP EAQVTFATPKKPGKEACSSIQV LNRGYRPPTFSYNTGVRRSARD YYPRRQRSISTNSPSRRWGKL	24208	54576	A	24346	880	1103	
GSGCAGLSL 24209 54577 A 24347 I 952 MITLRKLPLAVVAAGVMSAQ AMAVDFHGYARSGIGWTGSQG EQQCFQTTGAQSKYRLGMECE TYAELKLGGEVWKEGDKSFYF DTNVAYSVAQQNDWEATDPAF REANVQGKNLEWLPGSTIWA GKRFYQRHDVHMIDFYYWDIS GPGAGLENIDVGFGKLSLAATR SSEAGGSSFASHHTYDYSNEN ANDVFDVRLARWDSPGGPLTG FPPSALACTSGVSGGKGVPGPP EAQVTFATPKKPGKEACSSIQV LNRGYRPPTFSYNTGVRRSARD YYPRRQRSISTINSPSRWKGKL		1	1		1		
24209 54577 A 24347 I 952 MITLRKLPLAVAVAAGVMSAQ AMAVDFHGYARSGIGWTGSGG EQQCFQTTGAGSKYRLGNEGE TYAELKLGGEVWKEGŪKSFVF DTNVAYSVAQQNDWEATDPAF REANVOGKNLIEWLPOSTIWA GKRFYQRIĐVHMIDFYYWDIS GPGAGLENIDVGGKLSLAATR SSEAGGSSSFASNHIYDYS/NEN AND/VEDVRLAR WDSPGGPLTG FPPSALACTSGVSGGKGVFGPP EAQVTFATPKKPGKEACSSIQV LNRG YRPPTFSYNTGKRSARD YYPPRRQRSISTNSPSRR WKGKL	l	l			ŀ		AEESVAETRLCP*CRLYAEYDP
AMAVDFHOVARSGIGWTOSGG EQQCFQTTGAQSKYRLGNECE TYAELKLGGEVWKEGDKSFYF DTNVAYSVAQQNDWEATDPAF REANVQGKNLEWLPGSTIWA GKRYQRHDVHMIDFYYWDIS GPGAGLENIDVGFGKLSLAATR SSEAGGSSSFASNHIYDYSNEN ANDVFDVRLARWDSPGGPLTG FPPSALACTSGVSG(GKPGPFP EAQVTFATPKKPGKEACSSIQV LNRGYRPPTFSYNTGVRRARD YYPPRQRSISTNSPSRRWKGKL		1			ŀ		GSGCAGLSL
EQQFQTTGAQSKYRLGNECE TYAELKLQQEVWKEGDKSFYF DTNVAYSVAQQNDWEATDPAF REANVQGKKLLEWLPGSTIWA GKRFYQRHDVHMIDFYYWDIS GPGAGLENIDVGFGKLSLAATR SSEAGGSSFASNHIYDYSNEN ANDVFDVRLARWDSPGGPLTG FPPSALACTSGVSGQKGYPGPP EAQVTFATPKKPGKEACSSIQV LNRGYRPPTESYNTGVRRSARD YYPPRQRSISTNSPSRRWGKL	24209	54577	Α	24347	1	952	MITLRKLPLAVAVAAGVMSAQ
TYAELKL.GOEVWKEGDKSFYF DTNVAYSVAQQNDWEATDPAF REANVQGKNLIEWLPGSTIWA GKRFYQRHDVHMIDFYYWDIS GPGAGLENIDVGFGKLSLAATR SSEAGGSSSFASNHIYDYS/NEN ANDVFDVRLARWDSPGGPLTG FPPSALACTSGVSGGKGVFGPP EAQVTFATPKKPGKEACSSIQV LNRGYRPPTFSYNTGVRSARD YYPPRQRSISTNSPSRRWKGKL			1		l .		AMAVDFHGYARSGIGWTGSGG
DTNVAYSVAQQNDWEATDPAF REANVOGKNLIEWLPOSTIWA GKREYQRHDVHIMPFYYWDIS GPGAGLENIDVGFGKLSLAATR SSEAGGISSFASNHIYDYSINEN ANDVFDWILA RAWDSPIGGELTG FPPSALACTSGVSGQKGVPGPP EAQVITEATPKKPGKEACSSIQV LNRGYRPPTESYNTGVRRSARD YYPRRQRSISTNSPSRR WKGKL		1	1		l .		EOOCFOTTGAQSKYRLGNECE
DTNVAYSVAQQNDWEATDPAF REANVOGKNLIEWLPOSTIWA GKREYQRHDVHIMPFYYWDIS GPGAGLENIDVGFGKLSLAATR SSEAGGISSFASNHIYDYSINEN ANDVFDWILA RAWDSPIGGELTG FPPSALACTSGVSGQKGVPGPP EAQVITEATPKKPGKEACSSIQV LNRGYRPPTESYNTGVRRSARD YYPRRQRSISTNSPSRR WKGKL		1	l		l .		TYAELKLGOEVWKEGDKSFYF
REANVQGKNLIEWLPGSTIWA GKRFYQRHDVHMIDFYYWDIS GPGAGLENIDVGGKLSLAATR SSEAGGSSFASNHIYDYSNEN ANDAYEDVRLARWDSPGGPLTG FPPSALACTSGVSGQKGVPGPP EAQVTFATPKKPGKEACSSIQV LNRGYRPPTESYNTGVRRSARD YYPPRQRSISTNSPSRRWKGKL	1	l .	1		Ì		
GKRFYQRHDVHMIDFYYWDIS GPGAGLENIDVGFGKLSLAATR SSEAGGSSFASNHIYDYSNEN ANDVFDVRLARWDSPGGPLTG FPPSALACTSGVSGGKGVFGPP EAQVTFATPKKPGKEACSSIQV LNRGYRPPTFSYNTGVRRSARD YYPRRQRSISTNSPSRRWKGKL	ļ		1		ŀ		
GPGAGLENIDVGFGKLSLAATR SSEAGGSSFASNHIYDYSNEN ANDVFDVRLARWDSFGGHLFG FPPSALACTSGVSGQKGVPGPP EAQVTFATPKKPGKEACSSIQV LNRGYRPPTESYNTGVRRSARD YYPRRQRSISTNSPSRRWGKL							
SSEAGGSSSFASNHIYDYS/NEN AND/VEDVRLAR/VDSPGOPLTG FPPSALACTISGVSGGKOYOFPP EAQVIFATPKKPGKEACSSIQV LNRGYRPPTFSYNTGVRSARD YYPPRGRSISTNSPSRRWKGKL					i		
AND/VFDVRLARWDSPGGPLTG FPPSALACTSGVSGQKGVPGPP EAQVTFATPKKPGKEACSSIQV LNRGYRPPTESYNTGVRSSARD YYPRRQRSISTNSPSRRWKGKL		1	1	1	1	I	
FPPSALACTSGVSGQKGVPGPP EAQVTFATPKKPGKEACSSIQV LNRGVRPPTFSYTGVTRSABO YYPRRQRSISTNSPSRRWKGKL		1	1	1		I	
EAQVTFATPKKPGKEACSSIQV LNRGYRPPTFSYNTGVRRSARD YYPRRGRSISTNSPSRRWGGKL	1	1	1	1	ĺ	I	
LNRGYRPPTFSYNTGVRRSARD YYPRRQRSISTNSPSRRWKGKL		1	1	1	1	I	
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No.	
24210 54578 A 24348 805 1274 ASPATTRPVTKLA VGQRYWGSAASI ILASIFR/DRIASLE MS*AW_WGGP QFFSPARSASSHF IALGYRSESSSG RDYAHLEGLEPD C C C C C C C C C	
VGQRYWGSAASI ILASIFRDRIASLE MS*AWYWGCFF QFFSPARSASSHF IALGYRSESSEC RDYAHLHGLRFD C C C C C C C C C	
VGQRYWGSAASI ILASIFRDRIASLE MS*AWYWGCFF QFFSPARSASSHF IALGYRSESSEC RDYAHLHGLRFD C C C C C C C C C	MOVATEDEV
ILASIFR/DRIASLE MS*A WYWQCCF	
MS1AWYWQGCF	
OFFSPARSASSHE IALGYRSSESSIC RDYAHLHGLRFD C	
ALGYRSSESSCI RDVAHLHGLRED C C C C C C C C C	
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242 11 54579	HSSSKKKLCY
24212 54580 A 24350 I 696	
24213 54581 A 24351 I 735 MFTPDYRSVSRV MNQPDYMPDIRG HMIDFYYWDISG GFGKLSLAATRS: SNNIYDYTNETAH QMEINFGGTLELL RDNYRLGKGLSQ KFAYNINNNGHL MGDNWDMMYV NDHGJTKWWTA PHMYTGMEFGY QNIRTIYLAKPYQ QNIRTIYLA QNIRTIYLA QNIRTIYLA QNIRTIY QNIRTIYLA QNIRTIYLA QNIRTIY QNIRTIYLA QNIRTIY	
MNQPDVMPDIPG HMIDFYYVDISG GFGKLSLAATRS: SNNIVDYTNETAI QMEINPGGTLELL RDNYRLGKGLSQ KFAYNINNGHIM MGDNWDMMYV- NDHGYTKWWTA PIMNTGMKF/GYF QNIRTIYLAKPPQ	COCROVACE
HMIDFYYWDISG GFGKLSLAATRS: SNNI/DYTNETAI QMEINPGGTLELC RDNYRLGKGLSG KF4YNINNGH MGDNWDMMYV NDHGTKWWTA PIMNTGMKF/GYI QNIRTIYLAKPYQ	
GFGKLSLAATRS: SNNIYDYTNETAI QMEINFGGTLELC RDNYRLGKGLSQ KFAYNINNNGHI MGDNWDMMYV. NDHGTKWWTA PIMNTGMKF/GYI QNIRTIYLAKPPQ	
SNNIVDYTNETAL QMEINPGGTLELK RDNYRLGKGLSQ KFAYNINNGHI MGDNWDMMYV NDHGITKIWWTA PIMNTGMKFGYH QNIRTIYLAKPPQ	
QMEINPGGTLELG RDNYRLGKGLSQ KF4YNINNDH MGDNWDMMYV NDHGTKWWTA PIMNTGMKF/GY QNIRTIYLAKPYQ	
RDNYRLGKGLSQ KFAYNINNGHM MGDNWDMMYV NDHGITKIWWTA PIMNTGMKF/GYI QNIRTIYLAKPPQ	
KFA YNINNNGHM MGDWYDMMYV- NDHG/TK/WWTA PIIMTIGMKP/GYI QNIRTIYLAKP*Q	
MGDNWDMMYV NDHGTKIWWTA PIMNTGMKE/GYI QNIRTIYLAKP*Q	
NDHGTK\WWTA PIMNTGMKF/GYI QNIRTIYLAKPYQ	
PIMNTGMKF/GYI QNIRTIYLAKP*Q	
QNIRTIYLAKP*Q	
RFYQRHDVHMID	
GAGLENIDVGFG	
EAGGSSSFASNNI	
DVFDVRLAQMEI	
VDYGRANLRDNY	
SGVAFDNEKFAY	
RILDHGAISMGD	
MYQDIHWDNDH	
IRPDVQVDAHHE	
NPRARDKTFVQF	
24214 54582 A 24352 I 3414 MIVQPLAIIGSHFI	
PIAVYGRNEITAI	
ALRGTVSDVRKG	
IVAGNNDLSSRTE	
AASMEQLTATVO	
ASELAKNAATTA	
MTHTMQEIATSS	
GIAFQTNILALNA	AVEAARAGE
QGRGFAVVAGEV	RNLASRSAQ
AAKEIKGLIEESV	NRVQQGSKL
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